

VERIFICATION OF TRANSLATION

I, Shoji MIWA, a patent attorney of c/o Subaru Patent Office, Kojimachi Koyo Bldg., 10, Kojimachi 1-chome, Chiyoda-ku, Tokyo, Japan, hereby declare:

1. that I know well both the Japanese and English languages;
2. that the attached English translation is a true and correct translation of Japanese Patent Application No. 2000-321821 filed on October 20, 2000, priority of which is claimed in US Patent Application Serial No. 10/645,085 (Filing or 371 (c) Date: October 7, 2002), to the best of my knowledge and belief; and
3. that all statements made of my own knowledge are true and that all statements made on information and belief are believed to be true, and further that these statements are made with the knowledge that willful false statements and the like are punishable by fine or imprisonment, or both, under 18 USC 1001, and that such false statements may jeopardize the validity of the application or any patent issuing thereon.

Date: July 24, 2006

By: Shoji Miwa
Shoji MIWA

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**PATENT OFFICE
JAPANESE GOVERNMENT**

This is to certify that the annexed is a true copy of the following application as
filed with this Office.

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Japanese Patent Application Number: 2000-321821

Applicant(s): CHUGAI SEIYAKU KABUSHIKI KAISHA

Commissioner,
Patent Office

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[Identification of the Application]

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[Amendment 1]

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[Item to be amended] Inventor(s)

[Method of amendment] Change

[Contents of Amendment]

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[Note] The true inventors for the present application are **Naoshi FUKUSHIMA** and **Masayuki TSUCHIYA**, as declared in the written oath which is submitted herewith. However, in the application form, one of the true inventors, Naoshi FUKUSHIMA was mistakenly not listed as inventor, and Shinsuke UNO, who is not a joint inventor for this invention, was mistakenly listed as inventor. Accordingly, we hereby request to correct a section of inventor in the application form as identified above.

[Proof] Yes

[Document] Written Submission of Written Oath
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[Identification of the Application]

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[Contents of Supplement] Submission of Written Oath

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[Name of Deliverables] Written Oath 1

WRITTEN OATH

November 15, 2000

Patent Application No.: Japanese Patent Application No. 2000-321821
Title of the Invention: AGONIST ANTIBODIES

We, the undersigned, declare that the invention for the above-identified patent application has jointly invented by two inventors, **Naoshi FUKUSHIMA** and **Masayuki TSUCHIYA**, and **Shinsuke UNO** is not a joint inventor for the invention for the above-identified patent application.

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[Document] Specification

[Title of the Invention]

AGONIST ANTIBODIES

[Patent Claims]

5 [Claim 1] A modified antibody comprising two or more H chain V regions and two or more L chain V regions of monoclonal antibody and showing an agonist action by crosslinking a cell surface molecule(s).

10 [Claim 2] The modified antibody of claim 1, wherein the modified monoclonal antibody is a dimer of single chain Fv comprising an H chain V region and an L chain V region.

[Claim 3] The modified antibody of claim 1, wherein the modified antibody is a single chain polypeptide comprising two H chain V regions and two L chain V regions.

15 [Claim 4] The modified antibody of claim 2 or 3, wherein H chain V region and L chain V region are connected through a peptide linker comprises at least one amino acid.

[Claim 5] The modified antibody of any one of claims 1 to 4, wherein the modified antibody has been purified.

20 [Claim 6] The modified antibody of claim 1, wherein H chain V region and/or L chain V region is humanized H chain V region and/or L chain V region.

25 [Claim 7] The modified antibody of any one of claims 1 to 6, wherein the cell surface molecule is a hormone receptor or a cytokine receptor.

[Claim 8] The modified antibody of claim 7, wherein the cell surface molecule is selected from the group consisting of

erythropoietin (EPO) receptor, thrombopoietin (TPO) receptor, granulocyte colony stimulating factor (G-CSF) receptor, macrophage colony stimulating factor (M-CSF) receptor, granular macrophage colony stimulating factor (GM-CSF) receptor, tumor necrosis factor (TNF) receptor, interleukin-1 (IL-1) receptor, interleukin-2 (IL-2) receptor, interleukin-3 (IL-3) receptor, interleukin-4 (IL-4) receptor, interleukin-5 (IL-5) receptor, interleukin-6 (IL-6) receptor, interleukin-7 (IL-7) receptor, interleukin-9 (IL-9) receptor, interleukin-10 (IL-10) receptor, interleukin-11 (IL-11) receptor, interleukin-12 (IL-12) receptor, interleukin-13 (IL-13) receptor, interleukin-15 (IL-15) receptor, interferon-alpha (IFN-alpha) receptor, interferon-beta (IFN-beta) receptor, interferon-gamma (IFN-gamma) receptor, growth hormone (GH) receptor, insulin receptor, blood stem cell proliferation factor (SCF) receptor, vascular epidermal growth factor (VEGF) receptor, epidermal cell growth factor (EGF) receptor, nerve growth factor (NGF) receptor, fibroblast growth factor (FGF) receptor, platelet-derived growth factor (PDGF) receptor, transforming growth factor-beta (TGF-beta) receptor, leukocyte migration inhibitory factor (LIF) receptor, ciliary neurotrophic factor (CNTF) receptor, oncostatin M (OSM) receptor and Notch family receptor.

[Claim 9] The modified antibody of any one of claims 1 to 8, wherein the agonist action is induction of apoptosis, induction of cell proliferation and induction of cell

differentiation.

[Claim 10] A DNA which encodes the modified antibody of any one of claims 1 to 9.

5 [Claim 11] An animal cell which produces the modified antibody of any one of claims 1 to 9.

[Claim 12] A microorganism which produces the modified antibody of any one of claims 1 to 9.

[Claim 13] Use of the modified antibody of any one of claims 1 to 9 as an agonist.

10

[Detailed Description of the Invention]

[0001]

[Technical field to which the invention pertains]

15 This invention relates to modified antibodies containing two or more H chain V regions and two or more L chain V regions of a monoclonal antibody which show agonist activity by crosslinking a cell surface molecule(s). The modified antibodies have agonist activity of transducing a signal into cells by crosslinking a cell surface molecule(s)
20 which scan transduce a signal into cells and useful as a medicine for various purposes.

[0002]

[Prior art]

25 The inventors of this invention achieved the preparation of a specific monoclonal antibody using a splenic stromal cell line as a sensitizing antigen aiming at developing specific antibodies that can recognize the

aforementioned splenic stromal cells and the preparation of novel monoclonal antibodies that recognize mouse Integrin Associated Protein (mouse IAP) as an antigen. Then, the inventors also studied action of said antibodies using the recombinant body cells transfected with mouse IAP and found out that the monoclonal antibodies are capable of inducing apoptosis of myeloid cells. (JP-A 9-67499)
[0003]

Further, the inventors obtained hybridomas, MABL-1 (FERM BP-6100) and MABL-2 (FERM BP-6101), which can produce new monoclonal antibodies whose antigen is human Integrin Associated Protein (hereinafter referred to as human IAP; amino acid sequence and nucleotide sequence thereof are described in J. Cell Biol., 123, 485-496, 1993; see also Journal of Cell Science, 108, 3419-3425, 1995) and which are capable of inducing apoptosis of human nucleated blood cells (myeloid cell and lymphocyte) having said human IAP. The monoclonal antibodies produced by each hybridomas are referred to antibody MABL-1 and antibody MABL-2, respectively (WO99/12973). The monoclonal antibody recognizing IAP as an antigen induces apoptosis of nucleated blood cells having human IAP, but it also causes hemagglutination in vitro. It indicates that the administration of a large amount of the monoclonal antibody recognizing IAP as an antigen may result in a side effect such as hemagglutination.

[0004]

The inventors made intensive research for utilizing the monoclonal antibodies against human IAP as therapeutic agent of blood diseases and obtained single chain Fvs having the single chain Fv region capable of inducing apoptosis of nucleated blood cells having human IAP. (JP-A 11-63557)

[0005]

On the other hand modified antibodies, especially antibodies with lowered molecular size, for example, single chain Fvs were developed to improve permeability into tissues and tumors by lowering molecular size and to produce by a recombinant method. Recently the dimers of single chain Fvs, especially hetero-dimers are used for crosslinking cells. They are bispecific modified antibodies, whose typical example is hetero-dimers of single chain Fvs recognizing antigens of cancer cells and antigens of host cells like NK cells and neutrophils (Kipriyanov et al., Int. J. Cancer, 77, 9763-9772, 1998). They were produced by construction technique of single chain Fv as modified antibodies, which are more effective in treating cancers by inducing intercellular crosslinking. It has been thought that the intercellular crosslinking is induced by antibodies and their fragments (e.g. Fab fragment), bispecific modified antibodies and even dimers of single chain Fvs, which are monospecific.

[0006]

As antibodies capable of transducing a signal by crosslinking a cell surface molecule(s), there are known an antibody against EPO receptor involved in cell differentiation and proliferation (JP-A 2000-95800), an antibody against MuSK receptor (Xie et al., Nature Biotech. 15, 768-771, 1997) and others. However there have been no reports on modified antibodies with lowered molecular size.

[0007]

Noticing that antibody MABL-1, antibody MABL-2 and dimers derived from them induced apoptosis of cells having IAP, the inventors discovered that they crosslink (dimerize) IAP receptor on cell surface, thereby a signal is transduced into the cells and, as a result, apoptosis is induced. This suggests that monospecific single chain Fv dimers crosslink a cell surface molecule(s) (e.g. receptor) and transduce a signal like a ligand, thereby serving as an agonist.

[0008]

Focusing on the intercellular crosslinking, it was discovered that the above-mentioned single chain Fv dimers do not cause hemagglutination while the above-mentioned monoclonal antibodies do. The same result was also observed with single chain bivalent antibodies (single chain polypeptides containing two H chain V regions and two L chain V regions). This suggests that monoclonal antibodies may form intercellular crosslinking while modified antibodies like single chain Fv dimers and single chain

bivalent antibodies crosslink a cell surface molecule(s) but do not form intercellular crosslinking.

[0009]

Discovering that an antibody molecule (whole IgG) can be modified into single chain Fv dimers, single chain bivalent antibodies and the like which crosslink a cell surface molecule(s), thereby reducing side effects caused by intercellular crosslinking and providing new medicines inducing only desired effect on the cell, the inventors completed the invention. The modified antibodies have remarkably high activity compared with original monoclonal antibodies and improved permeability into tissues due to the characteristics of having lower molecular size compared with the original antibodies and of having no constant regions.

[0010]

[Problem to be solved by the invention]

Therefore, an object of this invention is to provide low molecular-size agonist modified antibodies which contain two or more H chain V regions and two or more L chain V regions of a monoclonal antibody and which combine with a cell surface molecule(s) and transduce a signal into cells, thereby can serve as an agonist.

[0011]

[Means to solve the problem]

This invention relates the modified antibodies which contain two or more H chain V regions and two or more

L chain V regions of a monoclonal antibody, and show an agonist activity by crosslinking a cell surface molecule(s).
[0012]

5 Preferable examples of the modified antibodies of
the invention are dimers of the single chain Fv which
contains one H chain V region and one L chain V region, or a
single chain polypeptide containing two H chain V regions
and two L chain V regions. The H chain V region and L chain
V region are preferably connected through a linker in the
10 modified antibodies.
[0013]

15 The above-mentioned single chain Fv dimer includes
a dimer by non-covalent bond, a dimer by a covalent bond
through a crosslinking radical and a dimer through a
crosslinking reagent (an antibody, an antibody fragment, or
a single chain Fv etc.). Conventional crosslinking radicals
used for crosslinking peptides can be used as the
crosslinking radicals to form the dimers. Examples are
disulfide crosslinking by cysteine residue, other
20 crosslinking radicals such as C₄ - C₁₀ alkylene (e.g.
tetramethylene, pentamethylene, hexamethylene,
heptamethylene and octamethylene, etc.) or C₄ - C₁₀
alkenylene (cis/trans -3-butenylene, cis/trans-2-pentenylene,
cis/trans-3-pentenylene, cis/trans-3-hexenylene, etc.).

25 Moreover, the crosslinking reagent which can
combine with a single chain Fv is, for example, an amino
acid sequence which can optionally be introduced into Fv,

for example, an antibody against FLAG sequence and the like or a fragment thereof, or a modified antibody originated from the antibody, for example, single chain Fv.

[0014]

5 The modified antibodies of this invention can be any things which contain L chain V region and H chain V region of monoclonal antibody (e.g. antibody MABL- 1, antibody MABL-2) and which specifically recognize the cell surface molecule(s), for example, a protein (a receptor or a protein involved in signal transduction), or a sugar chain
10 of the above-mentioned protein or of a cell membrane and crosslink said cell surface molecule(s), thereby transduce a signal into cells. Reconstructed polypeptides in which a part of amino acid sequence of V region has been altered are
15 included.

[0015]

 The present invention also relates to the humanization of the above-mentioned reconstructed polypeptides. The humanized reconstructed polypeptides
20 comprise a humanized H chain V region and/or a humanized L chain V region. Specifically, the humanized modified antibodies consist of the humanized L chain V region which comprises a framework region (FR) derived from an L chain V region of human monoclonal antibody and an CDR derived from
25 an L chain V region of mouse monoclonal antibody and/or the humanized H chain V region which comprises an FR derived from an H chain V region of human monoclonal antibody and a

CDR derived from an H chain V region of mouse monoclonal antibody. In this case, the amino acid sequence of FR or CDR may be partially altered, e.g. deleted, replaced or added.

[0016]

5 Furthermore, the present invention relates to polypeptides which comprise an L chain C region of human antibody and an L chain V region of the mouse monoclonal antibody, and/or an H chain C region of human antibody and an H chain V region of the mouse monoclonal antibody.

10 [0017]

 The present invention also relates to modified antibodies transducing a signal into cells by crosslinking a cell surface molecule(s), which comprise a CDR derived from a monoclonal antibody of other mammals than mouse (such
15 as human, rat, bovine, sheep, ape and the like), which is equivalent to said mouse CDR, or an H chain V region and an L chain V region containing the CDR. Such CDRs, H chain V regions and L chain V regions may include CDRs derived from a human monoclonal antibody prepared from, for example, a
20 transgenic mouse or the like, and H chain V regions and L chain V regions derived from a human monoclonal antibody containing the CDR.

[0018]

 The invention also relates to DNAs encoding the
25 various modified antibodies as mentioned above and genetic engineering techniques for the producing recombinant vectors comprising the DNAs.

[0019]

The invention also relates to host cells transformed with the recombinant vectors. Examples of host cells are animal cells such as human cells, mouse cells or the like and microorganisms such as E. coli, Bacillus subtilis, yeast or the like.

[0020]

The invention relates to a process for producing the modified antibodies, which comprises culturing the above-mentioned hosts and extracting the modified antibodies from the culture thereof.

[0021]

The present invention also relates to the use of the modified antibodies as an agonist. That is, it relates to the signal-transduction agonist which comprises as an active ingredient the modified antibody obtained as mentioned above. Since the modified antibodies used in the invention are those that crosslink the receptor on the cell surface and induce signal transduction, the receptor can be any receptor that is oligomerized, e.g. dimerized, by combining with the ligand and thereby transduce a signal into cells. The receptor includes hormone receptors and cytokine receptors. The hormone receptor includes, for example, estrogen receptor. The cytokine receptor and the like include hematopoietic factor receptor, lymphokine receptor, growth factor receptor, differentiation control factor receptor and the like. Examples of cytokine receptors

are erythropoietin (EPO) receptor, thrombopoietin (TPO) receptor, granulocyte colony stimulating factor (G-CSF) receptor, macrophage colony stimulating factor (M-CSF) receptor, granular macrophage colony stimulating factor (GM-CSF) receptor, tumor necrosis factor (TNF) receptor, interleukin-1 (IL-1) receptor, interleukin-2 (IL-2) receptor, interleukin-3 (IL-3) receptor, interleukin-4 (IL-4) receptor, interleukin-5 (IL-5) receptor, interleukin-6 (IL-6) receptor, interleukin-7 (IL-7) receptor, interleukin-9 (IL-9) receptor, interleukin-10 (IL-10) receptor, interleukin-11 (IL-11) receptor, interleukin-12 (IL-12) receptor, interleukin-13 (IL-13) receptor, interleukin-15 (IL-15) receptor, interferon-alpha (IFN-alpha) receptor, interferon-beta (IFN-beta) receptor, interferon-gamma (IFN-gamma) receptor, growth hormone (GH) receptor, insulin receptor, blood stem cell proliferation factor (SCF) receptor, vascular epidermal growth factor (VEGF) receptor, epidermal cell growth factor (EGF) receptor, nerve growth factor (NGF) receptor, fibroblast growth factor (FGF) receptor, platelet-derived growth factor (PDGF) receptor, transforming growth factor-beta (TGF-beta) receptor, leukocyte migration inhibitory factor (LIF) receptor, ciliary neurotrophic factor (CNTF) receptor, oncostatin M (OSM) receptor, Notch family receptor and the like.

Therefore, the pharmaceutical preparations containing the agonist modified antibody as an active ingredient are useful for as, for example, preventives and/or remedies for various

disease such as cancers, inflammation, hormone disorders and blood diseases.

[0022]

[Detailed Description of the Invention]

5 The modified antibodies of the present invention
comprise two or more H chain V regions and two or more L
chain V regions derived from monoclonal antibodies. The
structure of the reconstructed polypeptide may be a dimer of
single chain Fv comprising one H chain V region and one L
10 chain V region or a polypeptide comprising two H chain V
regions and two L chain V regions. In the reconstructed
polypeptide of the invention, the V regions of H chain and L
chain are preferably linked through a peptide linker which
consists of one or more amino acids. The resulting
15 reconstructed polypeptides contain variable regions of the
parent antibodies and retain the complementarity determining
region (CDR) thereof, and therefore bind to the antigen with
the same specificity as that of the parent monoclonal
antibodies.

20 [0023]

H chain V region

25 In the present invention, the H chain V region
derived from a monoclonal antibody recognizes a cell surface
molecule(s), for example, a protein (a receptor or a protein
involved in signal transduction) or a sugar chain of the
protein or on cell membrane and oligomerizes, for example,
dimerizes through crosslinking of said molecule, and thereby

serves as an agonist transducing a signal into the cells. The H chain V region of the invention includes H chain V regions derived from a mammal (e.g. human, mouse, rat, bovine, sheep, ape etc.) and partially modified H chain V regions thereof. More preferable is a humanized H chain V region containing FR of H chain V region of a human monoclonal antibody and CDR of H chain V region of a mouse monoclonal antibody. The H chain V region further can be an H chain V region derived from a human monoclonal antibody corresponding to the aforementioned H chain V region of mouse monoclonal antibody, which can be produced by recombination technique. The H chain V region of the invention may be a fragment of aforementioned H chain V region, which fragment preserves the antigen binding capacity.

[0024]

L chain V region

In the present invention, the L chain V region derived from the monoclonal antibody recognizes a cell surface molecule(s), for example, a protein (a receptor or a protein involved in signal transduction) or a sugar chain of the protein or on cell membrane and oligomerizes, for example, dimerizes through crosslinking of said molecule, and thereby serves as an agonist transducing a signal into the cells. The L chain V region of the invention includes L chain V regions derived from a mammal (e.g. human, mouse, rat, bovine, sheep, ape etc.) and partially modified L chain

V regions thereof. More preferable is a humanized L chain V region containing FR of L chain V region of human monoclonal antibody and CDR of L chain V region of mouse monoclonal antibodies. The L chain V regions further can be an L chain V region derived from human monoclonal antibody corresponding to the aforementioned L chain V region of mouse monoclonal antibody, which can be produced by recombination technique. The L chain V regions of the invention may be fragments of L chain V region, which fragments preserve the antigen binding capacity.

[0025]

Complementarity determining region (CDR)

Each V region of L chain and H chain forms an antigen-binding site. The variable region of the L and H chains is composed of comparatively conserved four common framework regions linked to three hypervariable regions or complementarity determining regions (CDR) (Kabat, E.A. et al., "Sequences of Protein of Immunological Interest", US Dept. Health and Human Services, 1983).

[0026]

Major portions in the four framework regions (FRs) form β -sheet structures and thus three CDRs form a loop. CDRs may form a part of the β -sheet structure in certain cases. The three CDRs are held sterically close position to each other by FR, which contributes to the formation of the antigen-binding site together with three CDRs.

[0027]

These CDRs can be identified by comparing the amino acid sequence of V region of the obtained antibody with known amino acid sequences of V regions of known antibodies according to the empirical rule in Kabat, E.A. et al., "Sequences of Protein of Immunological Interest".

[0028]

Single chain Fv

A single chain Fv is a polypeptide monomer comprising an H chain V region and an L chain V region linked each other which are derived from monoclonal antibodies. The resulting single chain Fvs contain variable regions of the parent monoclonal antibodies and preserve the complementarity determining region thereof, and therefore the single chain Fvs bind to the antigen by the same specificity as that of the parent monoclonal antibodies (JP-Appl. 11-63557). A part of the variable region and/or CDR of the single chain Fv of the invention or a part of the amino acid sequence thereof may be partially altered, for example deleted, replaced or added. The H chain V region and L chain V region composing the single chain Fv of the invention are mentioned before and may be linked directly or through a linker, preferably a peptide linker. The constitution of the single chain Fv may be [H chain V region]-[L chain V region] or [L chain V region]-[H chain V region]. In the present invention, it is possible to make the single chain Fv to form a dimer by non-covalent bond, by a covalent bond

through a crosslinking radical and a dimer through a crosslinking reagent capable of binding to single chain Fv (an antibody, an antibody fragment, or a single chain Fv etc.), from which the modified antibody of the invention can be formed.

[0029]

Single chain reconstructed polypeptide

The modified antibodies comprised of a single chain reconstructed polypeptide of the present invention comprising two or more H chain V regions and two or more L chain V regions, contain two or more H chain V regions and L chain V regions as mentioned above. Each region of the peptide should be arranged such that the single chain reconstructed polypeptide forms a specific steric structure, concretely mimicking a steric structure formed by the dimer of single chain Fv. For instance, the V regions are arranged in the order of the following manner:

[H chain V region]-[L chain V region]-[H chain V region]-[L chain V region]; or

[L chain V region]-[H chain V region]-[L chain V region]-[H chain V region],

wherein these regions are connected through a peptide linker, respectively.

[0030]

Linker

In this invention, the linkers for the connection between the H chain V region and the L chain V region may be

any peptide linker which can be introduced by the genetic engineering procedure or any linker chemically synthesized. For instance, linkers disclosed in literatures, e.g. Protein Engineering, 9(3), 299-305, 1996 may be used in the invention. If peptide linkers are required, the following are cited as example linkers:

Ser
Gly-Ser
Gly-Gly-Ser
Ser-Gly-Gly
Gly-Gly-Gly-Ser
Ser-Gly-Gly-Gly
Gly-Gly-Gly-Gly-Ser
Ser-Gly-Gly-Gly-Gly
Gly-Gly-Gly-Gly-Gly-Ser
Ser-Gly-Gly-Gly-Gly-Gly
Gly-Gly-Gly-Gly-Gly-Gly-Ser
Ser-Gly-Gly-Gly-Gly-Gly-Gly
(Gly-Gly-Gly-Gly-Ser)_n and
(Ser-Gly-Gly-Gly-Gly)_n

wherein n is an integer not less than one. Preferable length of the linker peptide is normally the range of 1 to 15 amino acids, preferably 2-12 amino acids, more preferably 3 - 10 amino acids. The method for introducing those linkers will be described in the explanation for DNA construction coding for modified antibodies of the invention.

[0031]

The chemically synthesized linkers, i.e. the chemical crosslinking agents, according to the invention can be any linkers conventionally employed for the linkage of peptides. Examples of the linkers may include N-hydroxy succinimide (NHS), disuccinimidyl suberate (DSS), bis(sulfosuccinimidyl)suberate (BS³), dithiobis(succinimidyl propionate) (DSP), dithiobis(sulfosuccinimidyl propionate) (DTSSP), ethylene glycolbis(succinimidyl succinate) (EGS), ethylene glycolbis(sulfosuccinimidyl succinate) (sulfo-EGS), disuccinimidyl tartrate (DST), disulfosuccinimidyl tartrate (sulfo-DST), bis[2-(succinimido oxycarbonyloxy)ethyl]sulfone (BSOCOES), bis[2-(sulfosuccinimido oxycarbonyloxy)ethyl]sulfone (sulfo-BSOCOES) or the like. These are commercially available.

[0032]

To form a dimer of the single chain Fv it is preferable for a linker to have a length capable of forming the dimer efficiently. Specifically, preferable is a linker composed of 2 to 12 amino acids, more preferably 3 to 10 amino acids or other linkers corresponding thereto.

[0033]

Preparation of modified antibodies

The modified antibodies can be produced by connecting, through the aforementioned linker, an H chain V region and an L chain V region derived from known or novel monoclonal antibodies specifically binding to a cell surface molecule(s). As examples of the single chain Fvs are cited

MABL1-scFv and MABL2-scFv comprising the H chain V region and the L chain V region derived from the antibody MABL-1 and the antibody MABL-2, respectively. As examples of the single chain polypeptides comprising two H chain V regions and two L chain V regions are cited MABL1-sc(Fv)₂ and MABL2-sc(Fv)₂ comprising the H chain V region and the L chain V region derived from the aforementioned antibodies.

[0034]

For the preparation of the polypeptide, a signal peptide may be attached to N-terminal of the polypeptide if the polypeptide is desired to be a secretory peptide. A well-known amino acid sequence useful for the purification of polypeptide such as the FLAG sequence may be attached for the efficient purification of the polypeptide. In this case a dimer can be formed by using anti-FLAG antibody.

[0035]

For the preparation of the modified antibody of the invention, it is necessary to obtain a DNA, i.e. a DNA encoding the single chain Fv or a DNA encoding reconstructed single chain polypeptide. These DNAs, especially for MABL1-scFv, MABL2-scFv, MABL1-sc(Fv)₂ and/or MABL2-sc(Fv)₂ are obtainable from the DNAs encoding the H chain V region and the L chain V region derived from said Fv. They are also obtainable by PCR method using those DNA as a template and amplifying the part of DNA contained therein encoding desired amino acid sequence with the aid of a pair of primers corresponding to both ends thereof.

[0036]

In the case where each V region having partially modified amino acid sequence is desired, the V regions in which one or some amino acids are modified, i.e. deleted, replaced or added can be obtained by a procedure known in the art using PCR. A part of the amino acid sequence in the V region is preferably modified by the PCR known in the art in order to prepare the reconstructed polypeptide which is sufficiently active against the specific antigen.

[0037]

For the determination of primers for the PCR amplification, it is necessary to decide the type of the H chain and L chain of the desired antibodies. In the case of antibody MABL-1 and the antibody MABL-2 it has been reported, however, that the antibody MABL-1 has κ type L chains and $\gamma 1$ type H chains and the antibody MABL-2 has κ type L chains and $\gamma 2a$ type H chains (JP-Appl. 11-63557). For the PCR amplification of the DNA encoding the H chain and L chain of the antibody MABL-1 and/or the antibody MABL-2, primers described in Jones, S.T. et al., Bio/Technology, 9, 88-89, 1991 may be employed.

[0038]

For the amplification of the L chain V regions of the antibody MABL-1 and the antibody MABL-2 using the polymerase chain reaction (PCR), 5'-end and 3'-end oligonucleotide primers are decided as aforementioned. In the same manner, 5'-end and 3'-end oligonucleotide primers

are decided for the amplification of the H chain V regions of the antibody MABL-1 and the antibody MABL-2.

[0039]

In embodiments of the invention, the 5'-end
5 primers which contain a sequence "GATC" providing the
restriction enzyme Hinf I recognition site at the
neighborhood of 5'-terminal thereof are used and the 3'-end
primers which contain a nucleotide sequence "CCCGGG"
providing the XmaI recognition site at the neighborhood of
10 5'-terminal thereof are used. Other restriction enzyme
recognition site may be used instead of these sites as long
as they are used for subcloning a desired DNA fragment into
a cloning vector.

[0040]

15 Specifically designed PCR primers are employed to
provide suitable nucleotide sequences at 5'-end and 3'-end
of the cDNAs encoding the V regions of the antibodies MABL-1
and MABL-2 so that the cDNAs are readily inserted into an
expression vector and appropriately function in the
20 expression vector (e.g. this invention devises to increase
transcription efficiency by inserting Kozak sequence). The
V regions of the antibodies MABL-1 and MABL-2 obtained by
amplifying by PCR using these primers are inserted into HEF
expression vector containing the desired human C region (see
25 WO92/19759). The cloned DNAs can be sequenced by using any
conventional process which comprises, for example, inserting

the DNAs into a suitable vector and then sequencing using the automatic DNA sequencer (Applied Biosystems).

[0041]

5 A linker such as a peptide linker can be introduced into the reconstructed polypeptide of the invention in the following manner. Primers which have partially complementary sequence with the primers for the H chain V regions and the L chain V regions as described above and which code for the N-terminal or the C-terminal of the linker are designed. Then, the PCR procedure can be carried out using these primers to prepare a DNA encoding the peptide linker having desired amino acid sequence and length. The DNAs encoding the H chain V region and the L chain V region can be connected through the resulting DNA to produce the DNA encoding the reconstructed polypeptide of the invention which has the desired peptide linker. Once the DNA encoding one of the reconstructed polypeptide is prepared, the DNAs encoding the reconstructed polypeptide with or without the desired peptide linker can readily be produced by designing various primers for the linker and then carrying out the PCR using the primers and the aforementioned DNA as a template.

[0042]

25 Each V region of the reconstructed polypeptide of the present invention can be humanized by using conventional techniques (e.g. Sato, K. et al., Cancer Res., 53, 1-6 (1993)). Once a DNA encoding a humanized Fv is prepared, a

humanized single chain Fv, a fragment of the humanized
single chain Fv, a humanized monoclonal antibody and a
fragment of the humanized monoclonal antibody can readily be
produced according to conventional methods. Preferably,
5 amino acid sequences of the V regions thereof may be
partially modified, if necessary.

[0043]

Furthermore, a DNA derived from other mammalian
origin, for example a DNA of human, can be produced in the
10 same manner as used to produce DNA encoding the H chain V
region and the L chain V region derived from mouse mentioned
in the above. The resulting DNA can be used to prepare an H
chain V region and an L chain V region of other mammal,
especially human origin, a single chain Fv derived from
15 human and a fragment thereof, and a monoclonal antibody of
human origin and a fragment thereof.

[0044]

As mentioned above, when the aimed DNAs encoding
the V regions of the reconstructed polypeptides and the V
20 regions of the humanized reconstructed polypeptides are
prepared, the expression vectors containing them and hosts
transformed with the vectors can be obtained according to
conventional methods. Further, the hosts can be cultured
according to a conventional method to produce the
25 reconstructed single chain Fv, the reconstructed humanized
single chain Fv, the humanized monoclonal antibodies and
fragments thereof. They can be isolated from cells or a

medium and can be purified into a homogeneous mass. For this purpose any isolation and purification methods conventionally used for proteins, e.g. chromatography, ultra-filtration, salting-out and dialysis, may be employed in combination, if necessary, without limitation thereto. [0045]

For the production of the modified antibodies of the present invention, any expression systems can be employed, for example, eukaryotic cells such as animal cells, e.g., established mammalian cell lines, filamentous fungi and yeast, and prokaryotic cells such as bacterial cells e.g., E. coli. Preferably, the reconstructed polypeptides of the invention are expressed in mammalian cells, for example COS7 cells or CHO cells.

[0046]

In these cases, conventional promoters useful for the expression in mammalian cells can be used. Preferably, human cytomegalovirus (HCMV) immediate early promoter is used. Expression vectors containing the HCMV promoter include HCMV-VH-HC γ 1, HCMV-VL-HCK and the like which are derived from pSV2neo (WO92/19759).

[0047]

Additionally, other promoters for gene expression in mammal cell which may be used in the invention include virus promoters derived from retrovirus, polyoma virus, adenovirus and simian virus 40 (SV40) and promoters derived from mammal such as human polypeptide-chain elongation

factor-1 α (HEF-1 α). SV40 promoter can easily be used according to the method of Mulligan, R.C., et al. (Nature 277, 108-114 (1979)) and HEF-1 α promoter can also be used according to the methods of Mizushima, S. et al. (Nucleic
5 Acids Research, 18, 5322 (1990)).

[0048]

Replication origin (ori) which can be used in the invention includes ori derived from SV40, polyoma virus, adenovirus, bovine papilloma virus (BPV) and the like. An
10 expression vector may contain, as a selection marker, phosphotransferase APH (3') II or I (neo) gene, thymidine kinase (TK) gene, E. coli xanthine-guanine phosphoribosyl transferase (Ecogpt) gene or dihydrofolate reductase (DHFR) gene.

15 [0049]

The antigen-binding activity of the reconstructed polypeptide as prepared above can be evaluated using the binding-inhibitory ability of original antibodies as an index. Concretely, the activity is evaluated in terms of the
20 absence or presence of concentration-dependent inhibition of the binding of said monoclonal antibody as an index.

[0050]

More in detail, animal cells transformed with an expression vector containing a DNA encoding the modified
25 antibody of the invention, e.g., COS7 cells or CHO cells, are cultured. The cultured cells and/or the supernatant of the medium or the modified antibody purified from them are

used to determine the binding to antigen. As a control is used a supernatant of the culture medium in which cells transformed only with the expression vector were cultured. In the case of an antigen, for example, the antibody MABL-1 and the antibody MABL-2, a test sample of the modified antibody of the invention or the supernatant of the control is added to mouse leukemia cell line, L1210 cells, expressing human IAP and then an assay such as the flow cytometry is carried out to evaluate the antigen-binding activity.

[0051]

In vitro evaluation of the signal transduction effect (apoptosis-inducing effect in the cases of the antibody MABL-1 and the antibody MABL-2) is performed in the following manner: A test sample of the above modified antibody is added to the cells which are expressing the antibody or cells into which the gene for the antibody has been introduced, and is evaluated by the change caused by the signal transduction, for example, whether cell death is induced in a manner specific to the human IAP-antigen.

[0052]

In vivo evaluation of the apoptosis-inducing effect, for example, in the case where the modified antibody recognizes human IAP (e.g. modified antibodies derived from the antibody MABL-1 and the antibody MABL-2) is carried out in the following manner: A mouse model of human myeloma is prepared. To the mice is intravenously administered the

monoclonal antibody or the reconstructed polypeptide of the invention, which induces apoptosis of nucleated blood cells having IAP. To mice of a control group is administered PBS alone. The induction of apoptosis is evaluated in terms of antitumor effect based on the change of human IgG content in serum of the mice and their survival time.

[0053]

The modified antibodies of the invention, which comprises two or more H chain V regions and two or more L chain V regions, may be a dimer of the single chain Fv comprising one H chain V region and one L chain V region, or a single chain polypeptide in which two or more H chain V regions and two or more L chain V regions are connected. It is considered that owing to such construction the peptide mimics three dimensional structure of the antigen binding site of the parent monoclonal antibody and therefore retains an excellent antigen-binding property.

[0054]

The modified antibodies of the invention has been remarkably lowered in the molecular size compared with antibody molecule (whole IgG), and, therefore, have superior permeability into tissues and tumors and higher activity than original monoclonal antibodies. Therefore, it is possible to transduce various signals into cells by properly selecting the original antibody which is modified. The pharmaceutical preparations containing them are useful for treating diseases curable by inducing signal transduction,

for example cancers, inflammation, hormone disorders as well as blood dyscrasia, for example, leukemia, malignant lymphoma, aplastic anemia, myelodysplasia syndrome and polycythemia vera. It is further expected that the antibody of the invention can be used as a contrast agent by RI-labeling. The effect can be enhanced by attaching to a RI-compound or a toxin.

[0055]

The present invention is illustrated by examples, which by no means restrict the scope of the invention, using monoclonal antibodies binding to human IAP (the antibody MABL-1 and the antibody MABL-2).

[0056]

[Examples]

For illustrating the production process of the reconstructed polypeptides of the invention, examples of producing single chain Fvs are shown below. Mouse antibodies against human IAP, MABL-1 and MABL-2 were used in the examples of producing the reconstructed polypeptides.

Hybridomas MABL-1 and MABL-2 producing them respectively were internationally deposited as FERM BP-6100 and FERM BP-6101 with the National Institute of Bioscience and Human Technology, Agency of Industrial Science and Technology, Minister of International Trade and Industry (1-3 Higasi 1-chome, Tsukuba-shi, Ibaraki-ken, Japan), an authorized depository for microorganisms, on September 11, 1997.

[0057]

Example 1 (Cloning of DNAs encoding V region of mouse monoclonal antibodies to human IAP)

DNAs encoding variable regions of the mouse monoclonal antibodies to human IAP, MABL-1 and MABL-2, were cloned as follows.

[0058]

1.1 Preparation of messenger RNA (mRNA)

mRNAs of the hybridomas MABL-1 and MABL-2 were obtained by using mRNA Purification Kit (Pharmacia Biotech).

[0059]

1.2 Synthesis of double-stranded cDNA

Double-stranded cDNA was synthesized from about 1 µg of the mRNA using Marathon cDNA Amplification Kit (CLONTECH) and an adapter was linked thereto.

[0060]

1.3 PCR Amplification of genes encoding variable regions of an antibody by

PCR was carried out using Thermal Cycler (PERKIN ELMER).

[0061]

(1) Amplification of a gene coding for L chain V region of MABL-1

Primers used for the PCR method are Adapter Primer-1 (CLONTECH) shown in SEQ ID No. 1, which hybridizes to a partial sequence of the adapter, and MKC (Mouse Kappa Constant) primer (Bio/Technology, 9, 88-89, 1991) shown in

SEQ ID No. 2, which hybridizes to the mouse kappa type L chain V region.

[0062]

50 μ l of the PCR solution contains 5 μ l of 10 \times PCR Buffer II, 2 mM $MgCl_2$, 0.16 mM dNTPs (dATP, dGTP, dCTP and dTTP), 2.5 units of a DNA polymerase, AmpliTaq Gold (PERKIN ELMER), 0.2 μ M of the adapter primer of SEQ ID No. 1, 0.2 μ M of the MKC primer of SEQ ID No. 2 and 0.1 μ g of the double-stranded cDNA derived from MABL-1. The solution was preheated at 94°C of the initial temperature for 9 minutes and then heated at 94°C for 1 minute, at 60°C for 1 minute and at 72°C for 1 minute 20 seconds in order. This temperature cycle was repeated 35 times and then the reaction mixture was further heated at 72°C for 10 minutes.

[0063]

(2) Amplification of cDNA encoding H chain V region of MABL-

1

The Adapter Primer-1 shown in SEQ ID No. 1 and MHC- γ 1 (Mouse Heavy Constant) primer (Bio/Technology, 9, 88-89, 1991) shown in SEQ ID No. 3 were used as primers for PCR.

[0064]

The amplification of cDNA was performed according to the method of the amplification of the L chain V region gene, which was described in Example 1.3-(1), except for using 0.2 μ M of the MHC- γ 1 primer instead of 0.2 μ M of the MKC primer.

[0065]

(3) Amplification of cDNA encoding L chain V region of MABL-

2

The Adapter Primer-1 of SEQ ID No. 1 and the MKC
5 primer of SEQ ID No. 2 were used as primers for PCR.

[0066]

The amplification of cDNA was carried out
according to the method of the amplification of the L chain
V region gene of MABL-1 which was described in Example 1.3-
10 (1), except for using 0.1 µg of the double-stranded cDNA
derived from MABL-2 instead of 0.1 µg of the double-stranded
cDNA from MABL-1.

[0067]

(4) Amplification of cDNA encoding H chain V region of MABL-

2

15 The Adapter Primer-1 of SEQ ID No. 1 and MHC-γ2a
primer (Bio/Technology, 9, 88-89, 1991) shown in SEQ ID No.
4 were used as primers for PCR.

[0068]

20 The amplification of cDNA was performed according
to the method of the amplification of the L chain V region
gene, which was described in Example 1.3-(3), except for
using 0.2 µM of the MHC-γ2a primer instead of 0.2 µM of the
MKC primer.

25 [0069]

1.4 Purification of PCR products

The DNA fragment amplified by PCR as described above was purified using the QIAquick PCR Purification Kit (QIAGEN) and dissolved in 10 mM Tris-HCl (pH 8.0) containing 1 mM EDTA.

[0070]

1.5 Ligation and Transformation

About 140 ng of the DNA fragment comprising the gene encoding the mouse kappa type L chain V region derived from MABL-1 as prepared above was ligated with 50 ng of pGEM-T Easy vector (Promega) in the reaction buffer comprising 30 mM Tris-HCl (pH 7.8), 10 mM MgCl₂, 10 mM dithiothreitol, 1 mM ATP and 3 units of T4 DNA Ligase (Promega) at 15°C for 3 hours.

[0071]

Then, 1 µl of the reaction mixture was added to 50 µl of E. coli DH5α competent cells (Toyobo Inc.) and the cells were stored on ice for 30 minutes, incubated at 42°C for 1 minute and stored on ice for 2 minutes again. 100 µl of SOC medium (GIBCO BRL) was added. The cells of E. coli were plated on LB (Molecular Cloning: A Laboratory Manual, Sambrook et al., Cold Spring Harbor Laboratory Press, 1989) agar medium containing 100 µg/ml of ampicillin (SIGMA) and cultured at 37°C overnight to obtain the transformant of E. coli.

[0072]

The transformant was cultured in 3 ml of LB medium containing 50 µg/ml of ampicillin at 37°C overnight and the

plasmid DNA was prepared from the culture using the QIAprep Spin Miniprep Kit (QIAGEN).

[0073]

5 The resulting plasmid comprising the gene encoding the mouse kappa type L chain V region derived from the hybridoma MABL-1 was designated as pGEM-M1L.

[0074]

10 According to the same manner as described above, a plasmid comprising the gene encoding the mouse H chain V region derived from the hybridoma MABL-1 was prepared from the purified DNA fragment and designated as pGEM-M1H.

[0075]

15 A plasmid comprising the gene encoding the mouse kappa type L chain V region derived from the hybridoma MABL-2 was prepared from the purified DNA fragment and designated as pGEM-M2L.

[0076]

20 A plasmid comprising the gene encoding the mouse H chain V region derived from the hybridoma MABL-2 was prepared from the purified DNA fragment and designated as pGEM-M2H.

[0077]

Example 2 (DNA Sequencing)

25 The nucleotide sequence of the cDNA encoding region in the aforementioned plasmids was determined using Auto DNA Sequencer (Applied Biosystem) and ABI PRISM Dye

Terminator Cycle Sequencing Ready Reaction Kit (Applied Biosystem) according to the manufacturer's protocol.

[0078]

5 The nucleotide sequence of the gene encoding the L chain V region from the mouse antibody MABL-1, which is included in the plasmid pGEM-M1L, is shown in SEQ ID No. 5.

[0079]

10 The nucleotide sequence of the gene encoding the H chain V region from the mouse antibody MABL-1, which is included in the plasmid pGEM-M1H, is shown in SEQ ID No. 6.

[0080]

The nucleotide sequence of the gene encoding the L chain V region from the mouse antibody MABL-2, which is included in the plasmid pGEM-M2L, is shown in SEQ ID No. 7.

15 [0081]

The nucleotide sequence of the gene encoding the H chain V region from the mouse antibody MABL-2, which is included in the plasmid pGEM-M2H, is shown in SEQ ID No. 8.

[0082]

20 Example 3 (Determination of CDR)

The V regions of L chain and H chain generally have a similarity in their structures and each four framework regions therein are linked by three hypervariable regions, i.e., complementarity determining regions (CDR). An amino acid sequence of the framework is relatively well conserved, while an amino acid sequence of CDR has extremely high variation (Kabat, E.A., et al., "Sequences of Proteins

of Immunological Interest", US Dept. Health and Human Services, 1983).

[0083]

On the basis of these facts, the amino acid sequences of the variable regions from the mouse monoclonal antibodies to human IAP were applied to the database of amino acid sequences of the antibodies made by Kabat et al. to investigate the homology. The CDR regions were determined based on the homology as shown in Table 1.

[0084]

Table 1

<u>Plasmid</u>	<u>SEQ ID No.</u>	<u>CDR(1)</u>	<u>CDR(2)</u>	<u>CDR(3)</u>
pGEM-M1L	5	43-58	74-80	113-121
pGEM-M1H	6	50-54	69-85	118-125
pGEM-M2L	7	43-58	74-80	113-121
pGEM-M2H	8	50-54	69-85	118-125

[0085]

Example 4 (Identification of Cloned cDNA Expression (Preparation of Chimera MABL-1 antibody and Chimera MABL-2 antibody.)

4.1 Preparation of vectors expressing chimera MABL-1 antibody

cDNA clones, pGEM-M1L and pGEM-M1H, encoding the V regions of the L chain and the H chain of the mouse antibody MABL-1, respectively, were modified by the PCR method and introduced into the HEF expression vector (W092/19759) to prepare vectors expressing chimera MABL-1 antibody.

[0086]

A forward primer MLS (SEQ ID No. 9) for the L chain V region and a forward primer MHS (SEQ ID No. 10) for the H chain V region were designed to hybridize to a DNA encoding the beginning of the leader sequence of each V region and to contain the Kozak consensus sequence (J. Mol. Biol., 196, 947-950, 1987) and HindIII restriction enzyme site. A reverse primer MLAS (SEQ ID No. 11) for the L chain V region and a reverse primer MHAS (SEQ ID No. 12) for the H chain V region were designed to hybridize to a DNA encoding the end of the J region and to contain the splice donor sequence and BamHI restriction enzyme site.

[0087]

100 µl of a PCR solution comprising 10 µl of 10 × PCR Buffer II, 2 mM MgCl₂, 0.16 mM dNTPs (dATP, dGTP, dCTP and dTTP), 5 units of DNA polymerase AmpliTaq Gold, 0.4 µM each of primers and 8 ng of the template DNA (pGEM-M1L or pGEM-M1H) was preheated at 94°C of the initial temperature for 9 minutes and then heated at 94°C for 1 minute, at 60°C for 1 minute and at 72°C for 1 minute 20 seconds in order. This temperature cycle was repeated 35 times and then the reaction mixture was further heated at 72°C for 10 minutes.

[0088]

The PCR product was purified using the QIAquick PCR Purification Kit (QIAGEN) and then digested with HindIII and BamHI. The product from the L chain V region was cloned into the HEF expression vector, HEF-κ and the product from

the H chain V region was cloned into the HEF expression vector, HEF- γ . After DNA sequencing, plasmids containing a DNA fragment with a correct DNA sequence are designated as HEF-M1L and HEF-M1H, respectively.

5 [0089]

4.2 Preparation of vectors expressing chimera MABL-2 antibodies

Modification and cloning of cDNA were performed in the same manner described in Example 4.1 except for using
10 pGEM-M2L and pGEM-M2H as template DNA instead of pGEM-M1L and pGEM-M1H. After DNA sequencing, plasmids containing DNA fragments with correct DNA sequences are designated as HEF-M2L and HEF-M2H, respectively.

[0090]

15 4.3 Transfection to COS7 cells

The aforementioned expression vectors were tested in COS7 cells to observe the transient expression of the chimera MABL-1 and MABL-2 antibodies.

[0091]

20 (1) Transfection with genes for the chimera MABL-1 antibody

COS7 cells were co-transformed with the HEF-M1L and HEF-M1H vectors by electroporation using the Gene Pulser apparatus (BioRad). Each DNA (10 μ g) and 0.8 ml of PBS with
25 1×10^7 cells/ml were added to a cuvette. The mixture was treated with pulse at 1.5 kV, 25 μ F of electric capacity.

[0092]

After the restoration for 10 minutes at a room temperature, the electroporated cells were transferred into DMEM culture medium (GIBCO BRL) containing 10% γ -globulin-free fetal bovine serum. After culturing for 72 hours, the supernatant was collected, centrifuged to remove cell fragments and recovered.

[0093]

(2) Transfection with genes coding for the chimera MABL-2 antibody

The co-transfection to COS7 cells with the genes coding for the chimera MABL-2 antibody was carried out in the same manner as described in Example 4.3-(1) except for using the HEF-M2L and HEF-M2H vectors instead of the HEF-M1L and HEF-M1H vectors. The supernatant was recovered in the same manner.

[0094]

4.4 Flow cytometry

Flow cytometry was performed using the aforementioned culture supernatant of COS7 cells to measure binding to the antigen. The culture supernatant of the COS7 cells expressing the chimera MABL-1 antibody or the COS7 cells expressing the chimera MABL-2 antibody, or human IgG antibody (SIGMA) as a control was added to 4×10^5 cells of mouse leukemia cell line L1210 expressing human IAP and incubated on ice. After washing, the FITC-labeled anti-human IgG antibody (Cappel) was added thereto. After incubating

and washing, the fluorescence intensity thereof was measured using the FACScan apparatus (BECTON DICKINSON).

[0095]

Since the chimera MABL-1 and MABL-2 antibodies
5 were specifically bound to L1210 cells expressing human IAP,
it is confirmed that these chimera antibodies have proper
structures of the V regions of the mouse monoclonal
antibodies MABL-1 and MABL-2, respectively (Figures 1-3).

[0096]

10 Example 5 (Preparation of reconstructed Single chain Fv
(scFv) of the antibody MABL-1 and antibody MABL-2)

5.1 Preparation of reconstructed single chain Fv of antibody
MABL-1

The reconstructed single chain Fv of antibody
15 MABL-1 was prepared as follows. The H chain V region and the
L chain V of antibody MABL-1, and a linker were respectively
amplified by the PCR method and were connected to produce
the reconstructed single chain Fv of antibody MABL-1. The
production method is illustrated in Figure 4. Six primers
20 (A-F) were employed for the production of the single chain
Fv of antibody MABL-1. Primers A, C and E have a sense
sequence and primers B, D and F have an antisense sequence.

[0097]

The forward primer VHS for the H chain V region
25 (Primer A, SEQ ID No. 13) was designed to hybridize to a DNA
encoding the N-terminal of the H chain V region and to
contain NcoI restriction enzyme recognition site. The

reverse primer VHAS for H chain V region (Primer B, SEQ ID No. 14) was designed to hybridize to a DNA coding the C-terminal of the H chain V region and to overlap with the linker.

5 [0098]

The forward primer LS for the linker (Primer C, SEQ ID No. 15) was designed to hybridize to a DNA encoding the N-terminal of the linker and to overlap with a DNA encoding the C-terminal of the H chain V region. The reverse primer LAS for the linker (Primer D, SEQ ID No. 16) was designed to hybridize to a DNA encoding the C-terminal of the linker and to overlap with a DNA encoding the N-terminal of the L chain V region.

[0099]

15 The forward primer VLS for the L chain V region (Primer E, SEQ ID No. 17) was designed to hybridize to a DNA encoding the C-terminal of the linker and to overlap with a DNA encoding the N-terminal of the L chain V region. The reverse primer VLAS-FLAG for L chain V region (Primer F, SEQ ID No. 18) was designed to hybridize to a DNA encoding the C-terminal of the L chain V region and to have a sequence encoding the FLAG peptide (Hopp. T. P. et al., Bio/Technology, 6, 1204-1210, 1988), two stop codons and EcoRI restriction enzyme recognition site.

20 [0100]

In the first PCR step, three reactions, A-B, C-D and E-F, were carried out and PCR products thereof were

purified. Three PCR products obtained from the first PCR step were assembled by their complementarity. Then, the primers A and F were added and the full length DNA encoding the reconstructed single chain Fv of antibody MABL-1 was amplified (Second PCR). In the first PCR, the plasmid pGEM-M1H encoding the H chain V region of antibody MABL-1 (see Example 2), a plasmid pSC-DP1 which comprises a DNA sequence encoding a linker region comprising: Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser (SEQ ID No. 19) (Huston, J.S., et al., Proc. Natl. Acad. Sci. USA, 85, 5879-5883, 1988) and the plasmid pGEM-M1L encoding the L chain V region of antibody MABL-1 (see Example 2) were employed as template, respectively.

[0101]

50 µl of the solution for the first PCR step comprises 5 µl of 10 × PCR Buffer II, 2 mM MgCl₂, 0.16 mM dNTPs, 2.5 units of DNA polymerase, AmpliTaq Gold (PERKIN ELMER), 0.4 µM each of primers and 5 ng each of template DNA. The PCR solution was preheated at 94°C of the initial temperature for 9 minutes and then heated at 94°C for 1 minute, at 65°C for 1 minute and at 72°C for 1 minute and 20 seconds in order. This temperature cycle was repeated 35 times and then the reaction mixture was further heated at 72°C for 7 minutes.

[0102]

The PCR products A-B (371bp), C-D (63bp) and E-F (384bp) were purified using the QIAquick PCR Purification

Kit (QIAGEN) and were assembled in the second PCR. In the second PCR, 98 μ l of a PCR solution comprising 120 ng of the first PCR product A-B, 20 ng of the PCR product C-D and 120 ng of the PCR product E-F, 10 μ l of 10 \times PCR Buffer II, 2mM MgCl₂, 0.16 mM dNTPs, 5 units of DNA polymerase AmpliTaq Gold (PERKIN ELMER) was preheated at 94°C of the initial temperature for 8 minutes and then heated at 94°C for 2 minutes, at 65°C for 2 minutes and at 72°C for 2 minutes in order. This temperature cycle was repeated twice and then 0.4 μ M each of primers A and F were added into the reaction, respectively. The mixture was preheated at 94°C of the initial temperature for 1 minutes and then heated at 94°C for 1 minute, at 65°C for 1 minute and at 72°C for 1 minute and 20 seconds in order. This temperature cycle was repeated 35 times and then the reaction mixture was further heated at 72°C for 7 minutes.

[0103]

A DNA fragment of 843 bp produced by the second PCR was purified and digested by NcoI and EcoRI. The resultant DNA fragment was cloned into pSCFVT7 vector. The expression vector pSCFVT7 contains a pelB signal sequence suitable for E. coli periplasmic expression system (Lei, S.P., et al., J. Bacteriology, 169, 4379-4383, 1987). After the DNA sequencing, the plasmid containing the DNA fragment encoding correct amino acid sequence of the reconstructed single chain Fv of antibody MABL-1 is designated as "pscM1" (see Figure 5). The nucleotide sequence and the amino acid

sequence of the reconstructed single chain Fv of antibody MABL-1 contained in the plasmid pscM1 are shown in SEQ ID No. 20.

[0104]

5 The pscM1 vector was modified by the PCR method to prepare a vector expressing the reconstructed single chain Fv of antibody MABL-1 in mammalian cells. The resultant DNA fragment was introduced into pCHO1 expression vector. This expression vector, pCHO1, was constructed by digesting DHFR-
10 ΔE-rvH-PM1-f (WO92/19759) with EcoRI and SmaI to eliminate the antibody gene and connecting the EcoRI-NotI-BamHI Adapter (Takara Shuzo) thereto.

[0105]

15 As a forward primer for PCR, Sal-VHS primer shown in SEQ ID No. 21 was designed to hybridize to a DNA encoding the N-terminal of the H chain V region and to contain SalI restriction enzyme recognition site. As a reverse primer for PCR, FRH1anti primer shown in SEQ ID No. 22 was designed to hybridize to a DNA encoding the end of the first framework
20 sequence.

[0106]

25 100 μl of PCR solution comprising 10 μl of 10 × PCR Buffer II, 2 mM MgCl₂, 0.16 mM dNTPs, 5 units of the DNA polymerase, AmpliTaq Gold, 0.4 μl M each of primer and 8 ng of the template DNA (pscM1) was preheated at 95°C of the initial temperature for 9 minutes and then heated at 95°C for 1 minute, at 60°C for 1 minute and at 72°C for 1 minute

and 20 seconds in order. This temperature cycle was repeated 35 times and then the reaction mixture was further heated at 72°C for 7 minutes.

[0107]

5 The PCR product was purified using the QIAquick
PCR Purification Kit (QIAGEN) and digested by SalI and MboII
to obtain a DNA fragment encoding the N-terminal of the
reconstructed single chain Fv of antibody MABL-1. The pscM1
vector was digested by MboII and EcoRI to obtain a DNA
10 fragment encoding the C-terminal of the reconstructed single
chain Fv of antibody MABL-1. The SalI-MboII DNA fragment and
the MboII-EcoRI DNA fragment were cloned into pCHO1-Igs
vector. After DNA sequencing, the plasmid comprising the
desired DNA sequence was designated as "pCHOM1" (see Figure
15 6). The expression vector, pCHO1-Igs, contains a mouse IgG1
signal sequence suitable for the secretion-expression system
in mammalian cells (Nature, 322, 323-327, 1988). The
nucleotide sequence and the amino acid sequence of the
reconstructed single chain Fv of antibody MABL-1 contained
20 in the plasmid pCHOM1 are shown in SEQ ID No. 23.

[0108]

5.2 Preparation of reconstructed single chain Fv of antibody MABL-2

25 The reconstructed single chain Fv of antibody
MABL-2 was prepared in accordance with the aforementioned
Example 5.1. Employed in the first PCR step were plasmid
pGEM-M2H encoding the H chain V region of MABL-2 (see

Example 2) instead of pGEM-M1H and plasmid pGEM-M2L encoding the L chain V region of MABL-2 (see Example 2) instead of pGEM-M1L, to obtain a plasmid pscM2 which comprises a DNA fragment encoding the desired amino acid sequence of the single chain Fv of antibody MABL-2. The nucleotide sequence and the amino acid sequence of the reconstructed single chain Fv of antibody MABL-2 contained in the plasmid pscM2 are shown in SEQ ID No. 24.

[0109]

The pscM2 vector was modified by the PCR method to prepare a vector, pCHOM2, for the expression in mammalian cells which contains the DNA fragment encoding the correct amino acid sequence of reconstructed the single chain Fv of antibody MABL-2. The nucleotide sequence and the amino acid sequence of the reconstructed single chain Fv of antibody MABL-2 contained in the plasmid pCHOM2 are shown in SEQ ID No. 25.

[0110]

5.3 Transfection to COS7 cells

The pCHOM2 vector was tested in COS7 cells to observe the transient expression of the reconstructed single chain Fv of antibody MABL-2.

[0111]

The COS7 cells were transformed with the pCHOM2 vector by electroporation using the Gene Pulser apparatus (BioRad). The DNA (10 µg) and 0.8 ml of PBS with 1×10^7

cells/ml were added to a cuvette. The mixture was treated with pulse at 1.5 kV, 25 μ F of electric capacity.

[0112]

After the restoration for 10 minutes at a room temperature, the electroporated cells were transferred into IMDM culture medium (GIBCO BRL) containing 10% fetal bovine serum. After culturing for 72 hours, the supernatant was collected, centrifuged to remove cell fragments and recovered.

[0113]

5.4 Detection of the reconstructed single chain Fv of antibody MABL-2 in culture supernatant of COS7 cells

The existence of the single chain Fv of antibody MABL-2 in the culture supernatant of COS7 cells which had been transfected with the pCHOM2 vector was confirmed by the Western Blotting method.

[0114]

The culture supernatant of COS7 cells transfected with the pCHOM2 vector and the culture supernatant of COS7 cells transfected with the pCHO1 as a control were subjected to SDS electrophoresis and transferred to REINFORCED NC membrane (Schleicher & Schuell). The membrane was blocked with 5% skim milk (Morinaga Nyu-gyo), washed with 0.05% Tween 20-PBS and mixed with an anti-FLAG antibody (SIGMA). The membrane was incubated at room temperature, washed and mixed with alkaline phosphatase-conjugated mouse IgG antibody (Zymed). After incubating and washing at room

temperature, the substrate solution (Kirkegaard Perry Laboratories) was added to develop color (Figure 7).

[0115]

5 A FLAG-peptide-specific protein was detected only in the culture supernatant of the pCHOM2 vector-introduced COS7 cells and thus it is confirmed that the reconstructed single chain Fv of antibody MABL-2 was secreted in this culture supernatant.

[0116]

10 5.5 Flow cytometry

Flow cytometry was performed using the aforementioned COS7 cells culture supernatant to measure the binding to the antigen. The culture supernatant of the COS7 cells expressing the reconstructed single chain Fv of antibody MABL-2 or the culture supernatant of COS7 cells transformed with pCHO1 vector as a control was added to 2×10^5 cells of the mouse leukemia cell line L1210 expressing human Integrin Associated Protein (IAP) or the cell line L1210 transformed with pCOS1 as a control. After incubating on ice and washing, the mouse anti-FLAG antibody (SIGMA) was added. Then the cells were incubated and washed. Then, the FITC labeled anti-mouse IgG antibody (BECTON DICKINSON) was added thereto and the cells were incubated and washed again. Subsequently, the fluorescence intensity was measured using the FACSscan apparatus (BECTON DICKINSON).

25 [0117]

Since the single chain Fv of antibody MABL-2 was specifically bound to L1210 cells expressing human IAP, it is confirmed that the reconstructed single chain Fv of antibody MABL-2 has an affinity to human Integrin Associated Protein (IAP) (see Figures 8-11).

[0118]

5.6 Competitive ELISA

The binding activity of the reconstructed single chain Fv of antibody MABL-2 was measured based on the inhibiting activity against the binding of mouse monoclonal antibodies to the antigen.

[0119]

The anti-FLAG antibody adjusted to 1 μ g/ml was added to each well on 96-well plate and incubated at 37°C for 2 hours. After washing, blocking was performed with 1% BSA-PBS. After incubating and washing at a room temperature, the culture supernatant of COS7 cells into which the secretion-type human IAP antigen gene (SEQ ID No. 26) had been introduced was diluted with PBS into twofold volume and added to each well. After incubating and washing at a room temperature, a mixture of 50 μ l of the biotinized MABL-2 antibody adjusted to 100 ng/ml and 50 μ l of sequentially diluted supernatant of the COS7 cells expressing the reconstructed single chain Fv of antibody MABL-2 were added into each well. After incubating and washing at a room temperature, the alkaline phosphatase-conjugated streptoavidin (Zymed) was added into each well. After

incubating and washing at a room temperature, the substrate solution (SIGMA) was added and absorbance of the reaction mixture in each well was measured at 405 nm.

[0120]

5 The results revealed that the reconstructed single chain Fv of antibody MABL-2 (MABL2-scFv) evidently inhibited concentration-dependently the binding of the mouse antibody MABL-2 to human IAP antigen in comparison with the culture supernatant of the PCHO1-introduced COS7 cells as a control
10 (Figure 12). Accordingly, it is suggested that the reconstructed single chain Fv of antibody MABL-2 has the correct structure of each of the V regions from the mouse monoclonal antibody MABL-2.

[0121]

15 5.7 Apoptosis-inducing Effect in vitro

 An apoptosis-inducing action of the reconstructed single chain Fv of antibody MABL-2 was examined by Annexin-V staining (Boehringer Mannheim) using the L1210 cells transfected with human IAP gene, the L1210 cells transfected
20 with the pCOS1 vector as a control and CCRF-CEM cells.

[0122]

 To each 1×10^5 cells of the above cells was added the culture supernatant of the COS7 cells expressing the reconstructed single chain Fv of antibody MABL-2 or the
25 culture supernatant of COS7 cells transfected with the pCHO1 vector as a control at 50% final concentration and the mixtures were cultured for 24 hours. Then, the Annexin-V

staining was performed and the fluorescence intensity was measured using the FACScan apparatus (BECTON DICKINSON).

[0123]

Results of the Annexin-V staining are shown in Figures 13-18, respectively. Dots in the left-lower region represent living cells and dots in the right-lower region represent cells at the early stage of apoptosis and dots in the right-upper region represent cells at the late stage of apoptosis. The results show that the reconstructed single chain Fv of antibody MABL-2 (MABL2-scFv) remarkably induced cell death of L1210 cells specific to human IAP antigen (Figures 13-16) and that the reconstructed single chain Fv also induced remarkable cell death of CCRF-CEM cells in comparison with the control (Figures 17-18).

[0124]

5.8 Expression of MABL-2 derived single chain Fv in CHO cells

CHO cells were transfected with the pCHOM2 vector to establish a CHO cell line which constantly expresses the single chain Fv (polypeptide) derived from the antibody MABL-2.

CHO cells were transformed with the pCHOM2 vector by the electroporation using the Gene Pulser apparatus (BioRad). A mixture of DNA (10 µg) and 0.7 ml of PBS with CHO cells (1×10^7 cells/ml) was added to a cuvette. The mixture was treated with pulse at 1.5 kV, 25 µF of electric capacity. After the restoration for 10 minutes at a room

temperature, the electroporated cells were transferred into nucleic acid free α -MEM medium (GIBCO BRL) containing 10% fetal bovine serum and cultured. The expression of desired protein in the resultant clones was confirmed by SDS-PAGE and a clone with a high expression level was selected as a cell line producing the single chain Fv derived from the antibody MABL-2. The cell line was cultured in serum-free medium CHO-S-SFM II (GIBCO BRL) containing 10 nM methotrexate (SIGMA). Then, the culture supernatant was collected, centrifuged to remove cell fragments and recovered.

[0125]

5.9 Purification of MABL-2 derived single chain Fv produced in CHO cells

The culture supernatant of the CHO cell line expressing the single chain Fv obtained in Example 5.8 was concentrated up to twenty times using a cartridge for the artificial dialysis (PAN130SF, ASAHI MEDICALS). The concentrated solution was stored at -20°C and thawed on purification.

Purification of the single chain Fv from the culture supernatant of the CHO cells was performed using three kinds of chromatography, i.e., Blue-sepharose, a hydroxyapatite and a gel filtration.

[0126]

(1) Blue-sepharose column chromatography

The concentrated supernatant was diluted to ten times with 20 mM acetate buffer (pH 6.0) and centrifuged to remove insoluble materials (10000 × rpm, 30 minutes). The supernatant was applied onto a Blue-sepharose column (20 ml) equilibrated with the same buffer. After washing the column with the same buffer, proteins adsorbed in the column were eluted by a stepwise gradient of NaCl in the same buffer, 0.1, 0.2, 0.3, 0.5 and up to 1.0 M. The pass-through fraction and each eluted fraction were analyzed by SDS-PAGE. The fractions in which the single chain Fv were confirmed (the fractions eluted at 0.1 to 0.3M NaCl) were pooled and concentrated up to approximately 20 times using CentriPrep-10 (AMICON).

[0127]

(2) Hydroxyapatite

The concentrated solution obtained in (1) was diluted to 10 times with 10 mM phosphate buffer (pH 7.0) and applied onto the hydroxyapatite column (20 ml, BIORAD). The column was washed with 60 ml of 10 mM phosphate buffer (pH 7.0). Then, proteins adsorbed in the column were eluted by a linear gradient of sodium phosphate buffer up to 200 mM (see Figure 19). The analysis of each fraction by SDS-PAGE confirmed the single chain Fv in fraction A and fraction B.

[0128]

(3) Gel filtration

Each of fractions A and B in (2) was separately concentrated with CentriPrep-10 and applied onto TSKgel

G3000SWG column (21.5 × 600 mm) equilibrated with 20 mM acetate buffer (pH 6.0) containing 0.15 M NaCl.

Chromatograms are shown in Figure 20. The analysis of the fractions by SDS-PAGE confirmed that both major peaks (AI and BI) are of desired single chain Fv. In the gel filtration analysis, the fraction A was eluted at 36 kDa of apparent molecular weight and the fraction B was eluted at 76 kDa. The purified single chain Fvs (AI, BI) were analyzed with 15% SDS polyacrylamide gel. Samples were treated in the absence or presence of a reductant and the electrophoresis was carried out in accordance with the Laemmli's method. Then the protein was stained with Coomassie Brilliant Blue. As shown in Figure 21, both AI and BI gave a single band at 35 kDa of apparent molecular weight, regardless of the absence or presence of the reductant. From the above, it is concluded that AI is a monomer of the single chain Fv and BI is a non-covalently bound dimer of the single chain Fv. The gel filtration analysis of the fractions AI and BI with TSKgel G3000SW column (7.5 × 60 mm) revealed that a peak of the monomer is detected only in the fraction AI and a peak of the dimer is detected only in the fraction BI (Figure 22).

[0129]

5.10 Construction of vector expressing single chain Fv derived from antibody MABL-2 in E. coli cell

The pscM2 vector was modified by the PCR method to prepare a vector effectively expressing the single chain Fv

from the antibody MABL-2 in E. coli cells. The resultant DNA fragment was introduced into pSCFVT7 expression vector.

[0130]

As a forward primer for PCR, Nde-VHSm02 primer shown in SEQ ID No. 27 was designed to hybridize to a DNA encoding the N-terminal of the H chain V region and to contain a start codon and NdeI restriction enzyme recognition site. As a reverse primer for PCR, VLAS primer shown in SEQ ID No. 28 was designed to hybridize to a DNA encoding the C-terminal of the L chain V region and to contain two stop codons and EcoRI restriction enzyme recognition site. The forward primer, Nde-VHSm02, comprises five point mutations in the part hybridizing to the DNA encoding the N-terminal of the H chain V region for the effective expression in E. coli.

[0131]

100 μ l of a PCR solution comprising 10 μ l of 10 x PCR Buffer #1, 1 mM $MgCl_2$, 0.2 mM dNTPs, 5 units of KOD DNA polymerase (all from TOYOB0), 1 μ M of each primer and 100 ng of a template DNA (pscM2) was heated at 98°C for 15 seconds, at 65°C for 2 seconds and at 74°C for 30 seconds in order. This temperature cycle was repeated 25 times.

[0132]

The PCR product was purified using the QIAquick PCR Purification Kit (QIAGEN) and digested by NdeI and EcoRI, and then the resulting DNA fragment was cloned into pSCFVT7 vector, from which pelB signal sequence had been

eliminated by the digestion with NdeI and EcoRI. After DNA sequencing, the resulting plasmid comprising a DNA fragment with the desired DNA sequence is designated as "pscM2DEm02" (see Figure 23). The nucleotide sequence and the amino acid sequence of the single chain Fv derived from the antibody MABL-2 contained in the plasmid pscM2DEm02 are shown in SEQ ID No. 29.

[0133]

5.11 Expression of single chain Fv derived from antibody

MABL-2 in E. coli cells

E. coli BL21(DE3)pLyss (STRATAGENE) was transformed with pscM2DEm02 vector to obtain a strain of E. coli expressing the single chain Fv derived from antibody MABL-2. The resulting clones were examined for the expression of the desired protein using SDS-PAGE, and a clone with a high expression level was selected as a strain producing the single chain Fv derived from antibody MABL-2.

[0134]

5.12 Purification of single chain Fv derived from antibody

MABL-2 produced in E.coli

A single colony of E. coli obtained by the transformation was cultured in 3 ml of LB medium at 28°C for 7 hours and then in 70 ml of LB medium at 28°C overnight. This pre-culture was transplanted to 7 L of LB medium and cultured at 28°C with stirring at 300 rpm using the Jar-fermenter. When an absorbance of the medium reached

O.D.=1.5, the bacteria were induced with 1 mM IPTG and then cultured for 3 hours.

[0135]

5 The culture medium was centrifuged (10000 × g, 10 minutes) and the precipitated bacteria were recovered. To the bacteria was added 50 mM Tris-HCl buffer (pH 8.0) containing 5 mM EDTA, 0.1 M NaCl and 1% Triton X-100 and the bacteria were disrupted by ultrasonication (out put: 4, duty cycle: 70%, 1 minute × 10 times). The suspension of
10 disrupted bacteria was centrifuged (12000 × g, 10 minutes) to precipitate inclusion body. Isolated inclusion body was mixed with 50 mM Tris-HCl buffer (pH 8.0) containing 5 mM EDTA, 0.1 M NaCl and 4% Triton X-100, treated by ultrasonication (out put: 4, duty cycle: 50%, 30 seconds × 2
15 times) again and centrifuged (12000 × g, 10 minutes) to isolate the desired protein as precipitate and to remove containment proteins included in the supernatant.

[0136]

20 The inclusion body comprising the desired protein was lysed in 50 mM Tris-HCl buffer (pH 8.0) containing 6 M Urea, 5 mM EDTA and 0.1 M NaCl and applied onto Sephacryl S-300 gel filtration column (5 × 90 cm, Amersham Pharmacia) equilibrated with 50 mM Tris-HCl buffer (pH 8.0) containing 4M Urea, 5 mM EDTA, 0.1 M NaCl and 10 mM mercaptoethanol at
25 a flow rate of 5 ml/minutes to remove associated single chain Fvs with high-molecular weight. The obtained fractions were analyzed with SDS-PAGE and the fractions with high

purity of the protein were diluted with the buffer used in the gel filtration up to $O.D_{280}=0.25$. Then, the fractions were dialyzed three times against 50 mM Tris-HCl buffer (pH 8.0) containing 5 mM EDTA, 0.1 M NaCl, 0.5 M Arg, 2 mM glutathione in the reduced form and 0.2 mM glutathione in the oxidized form in order for the protein to be refolded. Further, the fraction was dialyzed three times against 20 mM acetate buffer (pH 6.0) containing 0.15 M NaCl to exchange the buffer.

[0137]

The dialysate product was applied onto Superdex 200 pg gel filtration column (2.6 × 60 cm, Amersham Pharmacia) equilibrated with 20 mM acetate buffer (pH 6.0) containing 0.15 M NaCl to remove a small amount of high molecular weight protein which was intermolecularly crosslinked by S-S bonds. As shown in Figure 24, two peaks, major and sub peaks, were eluted after broad peaks which are expectedly attributed to an aggregate with a high molecular weight. The analysis by SDS-PAGE (see Figure 21) and the elution positions of the two peaks in the gel filtration analysis suggest that the major peak is of the monomer of the single chain Fv and the sub peak is of the non-covalently bound dimer of the single chain Fv.

[0138]

5.13 Apoptosis-inducing activity in vitro of single chain Fv derived from antibody MABL-2

An apoptosis-inducing action of the single chain Fv from antibody MABL-2 (MABL2-scFv) produced by the CHO cells and E. coli was examined according to two protocols by Annexin-V staining (Boehringer Mannheim) using the L1210 cells (hIAP/L1210) into which human IAP gene had been introduced.

In the first protocol sample antibodies at the final concentration of 3 µg/ml were added to 5×10^4 cells of hIAP/L1210 cell line and cultured for 24 hours. Sample antibodies, i.e., the monomer and the dimer of the single chain Fv of MABL-2 from the CHO cells obtained in Example 5.9, the monomer and the dimer of the single chain Fv of MABL-2 from E. coli obtained in Example 5.12, and the mouse IgG antibody as a control were analyzed. After culturing, the Annexin-V staining was carried out and the fluorescence intensity thereof was measured using the FACScan apparatus (BECTON DICKINSON).

In the second protocol sample antibodies at the final concentration of 3 µg/ml were added to 5×10^4 cells of hIAP/L1210 cell line, cultured for 2 hours and mixed with anti-FLAG antibody (SIGMA) at the final concentration of 15 µg/ml and further cultured for 22 hours. Sample antibodies of the monomer of the single chain Fv of MABL-2 from the CHO cells obtained in Example 5.9 and the mouse IgG antibody as a control were analyzed. After culturing, the Annexin-V staining was carried out and the fluorescence intensity thereof was measured using the FACScan apparatus.

Results of the analysis by the Annexin-V staining are shown in Figures 25-31. The results show that the dimers of the single chain Fv polypeptide of MABL-2 produced in the CHO cells and E. coli remarkably induced cell death (Figures 26, 27) in comparison with the control (Figure 25), while no apoptosis-inducing action was observed in the monomers of the single chain Fv polypeptide of MABL-2 produced in the CHO cells and E. coli (Figures 28, 29). When anti-FLAG antibody was used together, the monomer of the single chain Fv polypeptide derived from antibody MABL-2 produced in the CHO cells induced remarkably cell death (Figure 31) in comparison with the control (Figure 30).

[0139]

5.14 Antitumor effect of the monomer and the dimer of scFv/CHO polypeptide with a model mouse of human myeloma
(1) Quantitative measurement of human IgG in mouse serum

Measurement of human IgG (M protein) contained in mouse serum was carried out by the following ELISA. 100 μ L of goat anti-human IgG antibody (BIOSOURCE, Lot#7902) diluted to 1 μ g/mL with 0.1% bicarbonate buffer (pH 9.6) was added to each well on 96 wells plate (Nunc) and incubated at 4°C overnight so that the antibody was immobilized. After blocking, 100 μ L of the stepwisely diluted mouse serum or human IgG (CAPPEL, Lot#00915) as a standard was added to each well and incubated for 2 hours at a room temperature. After washing, 100 μ L of alkaline phosphatase-labeled anti-human IgG antibody (BIOSOURCE, Lot#6202) which had been

diluted to 5000 times was added, and incubation was carried out for 1 hour at a room temperature. After washing, a substrate solution was added. After incubation, absorbance at 405 nm was measured using the MICROPLATE READER Model 3550 (BioRad). The concentration of human IgG in the mouse serum was calculated based on the calibration curve obtained from the absorbance values of human IgG as the standard.

[0140]

(2) Preparation of antibodies for administration

The monomer and the dimer of the scFv/CHO polypeptide were respectively diluted to 0.4 mg/mL or 0.25 mg/mL with sterile filtered PBS(-) on the day of administration to prepare samples for the administration.

[0141]

(3) Preparation of a mouse model of human myeloma

A mouse model of human myeloma was prepared as follows. KPMM2 cells passaged in vivo (JP-Appl. 7-236475) by SCID mouse (Japan Clare) were suspended in RPMI1640 medium (GIBCO-BRL) containing 10% fetal bovine serum (GIBCO-BRL) and adjusted to 3×10^7 cells/mL. 200 μ L of the KPMM2 cell suspension (6×10^6 cells/mouse) was transplanted to the SCID mouse (male, 6 week-old) via caudal vein thereof, which had been subcutaneously injected with the asialo GM1 antibody (WAKO JUNYAKU, 1 vial dissolved in 5 mL) a day before the transplantation.

[0142]

(4) Administration of antibodies

The samples of the antibodies prepared in (2), the monomer (250 μ L) and the dimer (400 μ L), were administered to the model mice of human myeloma prepared in (3) via caudal vein thereof. The administration was started from three days after the transplantation of KPMM2 cells and was carried out twice a day for three days. As a control, 200 μ L of sterile filtered PBS(-) was likewise administered twice a day for three days via caudal vein. Each group consisted of seven mice.

[0143]

(5) Evaluation of antitumor effect of the monomer and the dimer of scFv/CHO polypeptide with the model mouse of human myeloma

The antitumor effect of the monomer and the dimer of scFv/CHO polypeptide with the model mice of human myeloma was evaluated in terms of the change of human IgG (M protein) concentration in the mouse serum and survival time of the mice. The change of human IgG concentration was determined by measuring it in the mouse serum collected at 24 days after the transplantation of KPMM2 cells by ELISA described in the above (1). The amount of serum human IgG (M protein) in the serum of the PBS(-)-administered group (control) increased to about 8500 μ g/mL, whereas the amount of human IgG of the scFv/CHO dimer-administered group was remarkably low, that is, as low as one-tenth or less than that of the control group. Thus, the results show that the dimer of scFv/CHO strongly inhibits the growth of the KPMM2

cells (Figure 32). As shown in Figure 33, a remarkable elongation of the survival time was observed in the scFv/CHO dimer-administered group in comparison with the PBS(-)-administered group.

5 [0144]

From the above, it is confirmed that the dimer of scFv/CHO has an antitumor effect for the human myeloma model mice. It is considered that the antitumor effect of the dimer of scFv/CHO, the reconstructed polypeptide of the invention, results from the apoptosis-inducing action of the reconstructed polypeptide.

[0145]

5.15 Hemagglutination Test

15 Hemagglutination test and determination of hemagglutination were carried out in accordance with "Immuno-Biochemical Investigation", Zoku-Seikagaku Jikken Koza, edited by the Biochemical Society of Japan, published by Tokyo Kagaku Dojin.

20 Blood was taken from a healthy donor using heparin-treated syringes and washed with PBS(-) three times, and then erythrocyte suspension with a final concentration of 2% in PBS(-) was prepared. Test samples were the antibody MABL-2, the monomer and the dimer of the single chain Fv polypeptide produced by the CHO cells, and the monomer and
25 the dimer of the single chain Fv polypeptide produced by E. coli, and the control was mouse IgG (ZYMED). For the investigation of the hemagglutination effect, round bottom

96-well plates available from Falcon were used. 50 μ L per well of the aforementioned antibody samples and 50 μ L of the 2% erythrocyte suspension were added and mixed in the well. After incubation for 2 hours at 37°C, the reaction mixtures were stored at 4°C overnight and the hemagglutination thereof was determined. As a control, 50 μ L per well of PBS(-) was used and the hemagglutination test was carried out in the same manner. The mouse IgG and antibody MABL-2 were employed at 0.01, 0.1, 1.0, 10.0 or 100.0 μ g/mL of the final concentration of the antibodies. The single chain Fvs were employed at 0.004, 0.04, 0.4, 4.0, 40.0 or 80.0 μ g/mL of the final concentration and further at 160.0 μ g/mL only in the case of the dimer of the polypeptide produced by E. coli. Results are shown in the Table 2. In the case of antibody MABL-2, the hemagglutination was observed at a concentration of more than 0.1 μ g/mL, whereas no hemagglutination was observed for both the monomer and the dimer of the single chain Fv.

[0146]

Table 2 Hemagglutination Test

	Control	0.01	0.1	1	10	100	$\mu\text{g/mL}$		
mIgG	-	-	-	-	-	-			
MABL-2 (intact)	-	-	+	+++	+++	++			
	Control	0.004	0.04	0.4	4	40	80	$\mu\text{g/mL}$	
scFv/CHO monomer	-	-	-	-	-	-	-		
scFv/CHO dimer	-	-	-	-	-	-	-		
	Control	0.004	0.04	0.4	4	40	80	160	$\mu\text{g/mL}$
scFv/E.coli monomer	-	-	-	-	-	-	-	-	
scFv/E.coli dimer	-	-	-	-	-	-	-	-	

[0147]

Example 6 Modified antibody $\text{sc}(\text{Fv})_2$ comprising two H chain
V regions and two L chain V regions and antibody MABL-2
scFvs having linkers with different length

6.1 Construction of plasmid expressing antibody MABL-2

$\text{sc}(\text{Fv})_2$

For the preparation of a plasmid expressing the
reconstructed polypeptide [$\text{sc}(\text{Fv})_2$] which comprises two H
chain V regions and two L chain V regions derived from the
antibody MABL-2, the aforementioned pCHOM2, which comprises
the DNA encoding scFv derived from the MABL-2 described
above, was modified by the PCR method as mentioned below and
the resulting DNA fragment was introduced into pCHOM2.

Primers employed for the PCR are EF1 primer (SEQ
ID NO: 30) as a sense primer, which is designed to hybridize
to a DNA encoding EF1 α , and an antisense primer (SEQ ID NO:

19), which is designed to hybridize to the DNA encoding C-terminal of the L chain V region and to contain a DNA sequence coding for a linker region, and VLLAS primer containing SalI restriction enzyme recognition site (SEQ ID NO 31).

[0148]

100 μ l of the PCR solution comprises 10 μ l of 10 \times PCR Buffer #1, 1 mM MgCl₂, 0.2 mM dNTPs (dATP, dGTP, dCTP and dTTP), 5 units of KOD DNA polymerase (Toyobo, Inc.), 1 μ M of each primer and 100 ng of the template DNA (pCHOM2). The PCR solution was heated at 94°C for 30 seconds, at 50°C for 30 seconds and at 74°C for 1 minute in order. This temperature cycle was repeated 30 times.

[0149]

The PCR product was purified using the QIAquick PCR Purification Kit (QIAGEN) and digested by SalI. The resultant DNA fragment was cloned into pBluescript KS⁺ vector (Toyobo, Inc.). After DNA sequencing, a plasmid comprising the desired DNA sequence was digested by SalI and the obtained DNA fragment was connected using Rapid DNA Ligation Kit(BOEHRINGER MANNHEIM) to pCHOM2 digested by SalI. After DNA sequencing, a plasmid comprising the desired DNA sequence is designated as "pCHOM2(Fv)₂" (see Figure 34). The nucleotide sequence and the amino acid sequence of the antibody MABL-2 sc(Fv)₂ region contained in the plasmid pCHOM2(Fv)₂ are shown in SEQ ID No. 32.

[0150]

6.2 Preparation of Plasmid expressing antibody MABL-2 scFvs having linkers with various length

The scFvs containing linkers with different length and the V regions which are designed in the order of [H chain]-[L chain] (hereinafter "HL") or [L chain]-[H chain] (hereinafter "LH") were prepared using, as a template, cDNAs encoding the H chain and the L chain derived from the MABL-2 as mentioned below.

[0151]

To construct HL type scFv the PCR procedure was carried out using pCHOM2(Fv)₂ as a template. In the PCR step, a pair of CFHL-F1 primer (SEQ ID NO: 33) and CFHL-R2 primer (SEQ ID NO: 34) or a pair of CFHL-F2 primer (SEQ ID NO: 35) and CFHL-R1 primer (SEQ ID NO: 36) and KOD polymerase were employed. The PCR procedure was carried out by repeating 30 times the temperature cycle consisting of 94°C for 30 seconds, 60°C for 30 seconds and 72°C for 1 minute in order to produce a cDNA for the H chain containing a leader sequence at 5'-end or a cDNA for the L chain containing FLAG sequence at 3'-end thereof. The resultant cDNAs for the H chain and the L chain were mixed and PCR was carried out by repeating 5 times the temperature cycle consisting of 94°C for 30 seconds, 60°C for 30 seconds and 72°C for 1 minute in order using the mixture as templates and the KOD polymerase. To the reaction mixture were added CFHL-F1 and CFHL-R1 primers and then the PCR reaction was performed by repeating 30 times of the aforementioned

temperature cycle to produce a cDNA for HL-0 type without a linker.

[0152]

To construct LH type scFv, the PCR reaction was
5 carried out using, as a template, pGEM-M2L and pGEM-M2H
which contain cDNAs encoding the L chain V region and the H
chain V region from the antibody MABL-2, respectively (see
JP- Appl. 11-63557). A pair of T7 primer (SEQ ID NO: 37) and
CFLH-R2 primer (SEQ ID NO: 38) or a pair of CFLH-F2 primer
10 (SEQ ID NO: 39) and CFLH-R1 (SEQ ID NO: 40) and the KOD
polymerase (Toyobo Inc.) were employed. The PCR reaction was
performed by repeating 30 times the temperature cycle
consisting of 94°C for 30 seconds, 60°C for 30 seconds and
72°C for 1 minute in sequential order to produce a cDNA of
15 an L chain containing a leader sequence at 5'-end or a cDNA
of an H chain containing FLAG sequence at 3'-end thereof.
The resultant cDNAs of the L chain and the H chain were
mixed and PCR was carried out using this mixture as
templates and the KOD polymerase by repeating 5 times the
20 temperature cycle consisting of 94°C for 30 seconds, 60°C
for 30 seconds and 72°C for 1 minute in order. To the
reaction mixture were added T7 and CFLH-R1 primers and the
reaction was performed by repeating 30 times of the
aforementioned temperature cycle. The reaction product was
25 used as a template and PCR was carried out using a pair of
CFLH-F4 primer (SEQ ID NO: 41) and CFLH-R1 primer by
repeating 30 times the temperature cycle consisting of 94°C

for 30 seconds, 60°C for 30 seconds and 72°C for 1 minute in order to produce a cDNA of LH-0 type without a linker.

[0153]

The resultant cDNAs of LH-0 and HL-0 types were
5 digested by EcoRI and BamHI restriction enzymes (Takara Shuzo) and the digested cDNAs were introduced into an expression plasmid INPEP4 for mammalian cells using Ligation High (Toyobo Inc.), respectively. Competent E. coli JM109 (Nippon Gene) was transformed with each plasmid and the
10 desired plasmids were isolated from the transformed E. coli using QIAGEN Plasmid Maxi Kit (QIAGEN). Thus plasmids pCF2LH-0 and pCF2HL-0 were prepared.

[0154]

To construct the expression plasmids of HL type
15 containing linkers with different size, pCF2HL-0, as a template, and CFHL-X3 (SEQ ID NO: 42), CFHL-X4 (SEQ ID NO: 43), CFHL-X5 (SEQ ID NO: 44), CFHL-X6 (SEQ ID NO: 45) or CFHL-X7 (SEQ ID NO: 46), as a sense primer, and BGH-1 (SEQ ID NO: 47) primer, as an antisense primer, which is
20 complementary with the vector sequence were employed. PCR reaction was carried out using the KOD polymerase by repeating 30 times the temperature cycle consisting of 94°C for 30 seconds, 60°C for 30 seconds and 72°C for 1 minute in order and the reaction products were digested by restriction
25 enzymes XhoI and BamHI (Takara Shuzo). The digested fragments were introduced between XhoI and BamHI sites in the pCF2HL-0 using Ligation High (Toyobo Inc.),

respectively. Competent E. coli JM109 was transformed with each plasmid and the desired plasmids were isolated from the transformed E. coli by using Qiagen Plasmid Maxi kit. Thus expression plasmids pCF2HL-3, pCF2HL-4, pCF2HL-5, pCF2HL-6 and pCF2HL-7 were prepared.

To construct expression plasmid for the transient expression in COS7 cells the plasmids pCF2HL-0, pCF2HL-3, pCF2HL-4, pCF2HL-5, pCF2HL-6 and pCF2HL-7 were digested by restriction enzymes EcoRI and BamHI (Takara Shuzo) and the resultant fragments of approximately 800 bp were purified with agarose gel electrophoresis. The obtained fragments were introduced between EcoRI and BamHI sites in an expression plasmid pCOS1 for the expression in mammalian cells by using Ligation High (Toyobo Inc.), respectively. Competent E. coli DH5 α (Toyobo Inc.) was transformed with each plasmid and the desired plasmids were isolated from the transformed E. coli using Qiagen Plasmid Maxi kit. Thus the expression plasmids CF2HL-0/pCOS1, CF2HL-3/pCOS1, CF2HL-4/pCOS1, CF2HL-5/pCOS1, CF2HL-6/pCOS1 and CF2HL-7/pCOS1 were prepared.

As a typical example of these plasmids, the construction of the plasmid CF2HL-0/pCOS1 is illustrated in Figure 35 and the nucleotide sequence and the amino acid sequence of MABL2-scFv <HL-0> contained in the plasmid are shown in SEQ ID No. 48. Nucleotide sequences and amino acid sequences of the linker regions in these plasmids are also shown in Figure 36.

[0155]

To construct the expression plasmids of LH type containing linkers with different size, pCF2LH-0, as a template, and CFLH-X3 (SEQ ID NO: 49), CFLH-X4 (SEQ ID NO: 50), CFLH-X5 (SEQ ID NO: 51), CFLH-X6 (SEQ ID NO: 52) or CFLH-X7 (SEQ ID NO: 53), as a sense primer, and BGH-1 primer, as an antisense primer, which is complementary with the vector sequence were employed. PCR reaction was carried out using the KOD polymerase by repeating 30 times the temperature cycle consisting of 94°C for 30 seconds, 60°C for 30 seconds and 72°C for 1 minute in order and the reaction products were digested by restriction enzymes XhoI and BamHI. The digested fragments were introduced into the pCF2LH-0 between XhoI and BamHI sites using Ligation High, respectively. Competent E. coli DH5α (Toyobo Inc.) was transformed with each plasmid and the desired plasmids were isolated from the transformed E. coli using Qiagen Plasmid Maxi kit. Thus expression plasmids pCF2LH-3, pCF2LH-4, pCF2LH-5, pCF2LH-6 and pCF2LH-7 were prepared.

To construct expression plasmid for the transient expression in COS7 cells the plasmids pCF2LH-0, pCF2LH-3, pCF2LH-4, pCF2LH-5, pCF2LH-6 and pCF2LH-7 were digested by restriction enzymes EcoRI and BamHI (Takara Shuzo) and the resultant fragments of approximately 800 bp were purified with agarose gel electrophoresis. The obtained fragments were introduced between XhoI and BamHI sites in an expression plasmid pCOS1 for the expression in mammalian

cells by using the Ligation High, respectively. Competent E. coli DH5 α (Toyobo Inc.) was transformed with each plasmid and the desired plasmids were isolated from the transformed E. coli using the Qiagen Plasmid Maxi kit. Consequently, the expression plasmids CF2LH-0/pCOS1, CF2LH-3/pCOS1, CF2LH-4/pCOS1, CF2LH-5/pCOS1, CF2LH-6/pCOS1 and CF2LH-7/pCOS1 were prepared.

As a typical example of these plasmids, the construction of the plasmid CF2LH-0/pCOS1 is illustrated in Figure 37 and the nucleotide sequence and the amino acid sequence of MABL2-scFv <LH-0> contained in the plasmid are shown in SEQ ID No. 54. Nucleotide sequences and amino acid sequences of the linker regions in these plasmids are also shown in Figure 38.

[0156]

6.3 Expression of scFvs and sc(Fv)₂ in COS7 cells

(1) Preparation of culture supernatant using serum-containing culture medium

The HL type and LH type of scFvs and sc(Fv)₂ were transiently expressed in COS7 cells (JCRB9127, Japan Health Sciences Foundation). COS7 cells were subcultured in DMEM media (GIBCO BRL) containing 10% fetal bovine serum (HyClone) at 37°C in carbon dioxide atmosphere incubator.

[0157]

The COS7 cells were transfected with CF2HL-0, 3 ~ 7/pCOS1, or CF2LH-0, 3 ~ 7/pCOS1 prepared in Example 6.2 or pCHOM2(Fv)₂ vectors by electroporation using the Gene Pulser

apparatus (BioRad). The DNA (10 μ g) and 0.25 ml of 2×10^7 cells/ml in DMEM culture medium containing 10% FBS and 5 mM BES (SIGMA) were added to a cuvette. After standing for 10 minutes the mixtures were treated with pulse at 0.17kV, 5 950 μ F of electric capacity. After the restoration for 10 minutes at room temperature, the electroporated cells were transferred into the DMEM culture medium (10%FBS) in 75 cm³ flask. After culturing for 72 hours, the culture supernatant was collected and centrifuged to remove cell fragments. The 10 culture supernatant was subjected to the filtration using 0.22 μ m bottle top filter (FALCON) to obtain the culture supernatant (hereinafter "CM").

[0158]

(2) Preparation of culture supernatant using serum-free 15 culture medium

Cells transfected in the same manner as (1) were transferred to the DMEM medium (10% FBS) in 75 cm³ flask and cultured overnight. After the culture, the supernatant was discarded and the cells were washed with PBS and then added 20 to CHO-S-SFM II medium (GIBCO BRL). After culturing for 72 hours, the culture supernatant was collected, centrifuged to remove cell fragments and filtered using 0.22 μ m bottle top filter (FALCON) to obtain CM.

[0159]

25 6.4 Detection of scFvs and sc(Fv)₂ in CM of COS7

The various MABL2-scFVs and sc(Fv)₂ in CM of COS7 prepared in the aforementioned Example 6.3 (2) were detected by Western Blotting method.

Each CM of COS7 was subjected to SDS-PAGE electrophoresis and transferred to REINFORCED NC membrane (Schleicher & Schuell). The membrane was blocked with 5% skim milk (Morinaga Nyu-gyo) and washed with TBS. Then an anti-FLAG antibody (SIGMA) was added thereto. The membrane was incubated at room temperature and washed. A peroxidase labeled mouse IgG antibody (Jackson Immuno Research) was added. After incubating and washing at room temperature, the substrate solution (Kirkegaard Perry Laboratories) was added to develop color (Figure 39).

[0160]

6.5 Flow cytometry

Flow cytometry was performed using the culture supernatants of COS7 cells prepared in Example 6.3 (1) to measure the binding of the MABL2-scFVs and sc(Fv)₂ to human Integrin Associated Protein (IAP) antigen. The culture supernatants to be tested or a culture supernatant of COS7 cells as a control was added to 2×10^5 cells of the mouse leukemia cell line L1210 expressing human IAP. After incubating on ice and washing, 10 µg/mL of the mouse anti-FLAG antibody (SIGMA) was added and then the cells were incubated and washed. Then, the FITC labeled anti-mouse IgG antibody (BECTON DICKINSON) was added thereto and the cells

were incubated and washed again. The fluorescence intensity was measured using the FACScan apparatus (BECTON DICKINSON). [0161]

5 The results of the flow cytometry show that the MABL2-scFvs having linkers with different length and the sc(Fv)₂ in the culture supernatants of COS7 have high affinity to human IAP (see Figure 40).

[0162]

6.6 Apoptosis-inducing Effect in vitro

10 An apoptosis-inducing action of the culture supernatants of COS7 prepared in Example 6.3 (1) was examined by Annexin-V staining (Boehringer Mannheim) using the L1210 cells transfected with human IAP gene (hIAP/L1210).

15 To 5×10^4 cells of the hIAP/L1210 cells were added the culture supernatants of COS7 cells transfected with each vectors or a culture supernatant of COS7 cells as a control at 10% of the final concentration and the mixtures were cultured for 24 hours. Then, the Annexin-V/PI staining
20 was performed and the fluorescence intensity was measured using the FACScan apparatus (BECTON DICKINSON). The results revealed that scFvs <HL3, 4, 6, 7, LH3, 4, 6, 7> and sc(Fv)₂ in CM of COS7 induced remarkable cell death of hIAP/L1210 cells. These results are shown in Figure 41.

25 [0163]

6.7 Construction of vectors for the expression of scFvs and sc(Fv)₂ in CHO cells

To isolate and purify MABL2-scFvs and sc(Fv)₂ from culture supernatant, the expression vectors for expressing in CHO cells were constructed as below.

The EcoRI-BamHI fragments of pCF2HL-0, 3 ~ 7, and pCF2LH-0, 3 ~ 7 prepared in Example 6.2 were introduced between EcoRI and BamHI sites in an expression vector pCHO1 for CHO cells using the Ligation High. Competent E. coli DH5α was transformed with them. The plasmids were isolated from the transformed E. coli using QIAGEN Plasmid Midi kit (QIAGEN) to prepare expression plasmids pCHOM2HL-0, 3 ~ 7, and pCHOM2LH-0, 3 ~ 7.

[0164]

6.8 Production of CHO cells expressing MABL2-scFvs <HL-0, 3 ~ 7>, MABL2-scFvs <LH-0, 3 ~ 7> and sc(Fv)₂ and preparation of the culture supernatants thereof

CHO cells were transformed with each of the expression plasmids pCHOM2HL-0, 3 ~ 7, and pCHOM2LH-0, 3 ~ 7, constructed in Example 6.7 and pCHOM2(Fv)₂ vector to prepare the CHO cells constantly expressing each reconstructed polypeptide. As a typical example thereof, the production of the CHO cells constantly expressing MABL2-scFv <HL-5> or sc(Fv)₂ is illustrated as follows.

[0165]

The expression plasmids pCHOM2HL-5 and pCHOM2(Fv)₂ were linearized by digesting with a restriction enzyme PvuI and subjected to transfection to CHO cells by electroporation using Gene Pulser apparatus (BioRad). The

DNA (10 µg) and 0.75 ml of PBS with 1×10^7 cells/ml were added to a cuvette and treated with pulse at 1.5 kV, 25 µF of electric capacity. After the restoration for 10 minutes at room temperature, the electroporated cells were

5 transferred into nucleic acid-containing α-MEM culture medium (GIBCO BRL) containing 10% fetal bovine serum and cultured. After culturing overnight, the supernatant was discarded. The cells were washed with PBS and added to nucleic acid-free α-MEM culture medium (GIBCO BRL)
10 containing 10% fetal bovine serum. After culturing for two weeks, the cells were cultured in a medium containing 10 nM (final concentration) methotrexate (SIGMA), then 50 nM and 100 nM methotrexate. The resultant cells were cultured in serum-free CHO-S-SFM II medium (GIBCO BRL) in a roller
15 bottle. The culture supernatant was collected, centrifuged to remove cell fragments and filtered using a filter with 0.22 µm of pore size to obtain CM, respectively.

According to the above, CHO cells which constantly express MABL2-scFvs <HL-0, -3, -4, -6, -7> and <LH-0, -3, -
20 4, -5, -6, -7> and CMs thereof were obtained.

[0166]

6.9 Purification of dimer of MABL2-scFv <HL-5> and sc(Fv)₂

According to Example 5.9, CMs prepared in Example 6.8 were concentrated and the MABL2-scFv <HL-5> and the
25 sc(Fv)₂ were purified using three types of chromatography methods, Blue-sepharose, hydroxyapatite and gel filtration.

[0167]

6.10 Evaluation of the binding activity of purified dimer of scFv <HL-5> and sc(Fv)₂ against antigen

Flow cytometry was performed using the purified dimer of MABL2-scFv <HL-5> and the purified sc(Fv)₂ in order to evaluate the binding to human Integrin Associated Protein (IAP) antigen. 10µg/ml of the purified dimer of MABL2-scFv <HL-5>, the purified sc(Fv)₂, the antibody MABL-2 as a positive control or a mouse IgG (Zymed) as a negative control was added to 2×10^5 cells of the mouse leukemia cell line L1210 expressing human IAP (hIAP/L1210) or the cell line L1210 transformed with pCOS1 (pCOS1/L1210) as a control. After incubating on ice and washing, 10µg/mL of the mouse anti-FLAG antibody (SIGMA) was added and then the cells were incubated and washed. FITC labeled anti-mouse IgG antibody (BECTON DICKINSON) was added thereto and the cells were incubated and washed again. Then the fluorescence intensity was measured using the FACScan apparatus (BECTON DICKINSON).

[0168]

Since the purified dimer of MABL2-scFv <HL-5> and the purified sc(Fv)₂ were specifically bound to hIAP/L1210 cells, it is confirmed that the dimer of scFv <HL-5> and the sc(Fv)₂ have high affinity to human IAP (see Figure 42).

[0169]

6.11 Apoptosis-inducing activity in vitro of purified dimer of scFv <HL-5> and sc(Fv)₂

An apoptosis-inducing action of the purified dimer of MABL2-scFv <HL-5> and the purified sc(Fv)₂ were examined by Annexin-V staining (Boehringer Mannheim) using the L1210 cells (hIAP/L1210) in which human IAP gene had been introduced and cells of human leukemic cell line CCRF-CEM. [0170]

Different concentrations of the purified dimer of MABL2-scFv <HL-5>, the purified MABL2-sc(Fv)₂, the antibody MABL-2 as a positive control or a mouse IgG as a negative control were added to 5×10^4 cells of hIAP/L1210 cell line or 1×10^5 cells of CCRF-CEM cell line. After culturing for 24 hours, the Annexin-V staining was carried out and the fluorescence intensity thereof was measured using the FACScan apparatus (BECTON DICKINSON). As a result the dimer of MABL2-scFv <HL-5> and the MABL2-sc(Fv)₂ remarkably induced cell death of hIAP/L1210 and CCRF-CEM in concentration-dependent manner (see Figure 43). [0171]

6.12 Hemagglutination Test of the purified dimer of scFv <HL-5> and the sc(Fv)₂

Hemagglutination test was carried out using different concentrations of the purified dimer of scFv <HL-5> and the purified sc(Fv)₂ in accordance with Example 5.15.

The hemagglutination was observed with the antibody MABL-2 as a positive control, whereas no hemagglutination was observed with both the single chain antibody MABL2-sc(Fv)₂ and the MABL2-scFv <HL-5>. Further,

there was no substantial difference in the hemagglutination between two buffers employed with the antibody MABL-2. These results are shown in Table 3.

[0172]

Hemagglutination Test

TABLE 3

Diluent : PBS

(μ g/ml)

	cont	28.9	14.45	7.225	3.6125	1.8063	0.9031	0.4516	0.2258	0.1129	0.0564	0.0282	0.0141	0.0071	0.0035	0.0018
MABL2-sc(Fv) ₂	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	cont	28.0	14.0	7.0	3.5	1.75	0.875	0.4375	0.2188	0.1094	0.0547	0.0273	0.0137	0.0068	0.0034	0.0017
MABL2-sc(Fv)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<HL5>	cont	80	40	20	10	5	2.5	1.25	0.625	0.3125	0.1563	0.0781	0.0391	0.0195	0.0098	0.0049
MABL2(intact)	-	+	+	+	+	+	+	+	+	+	±	-	-	-	-	-
mlgG	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Diluent : Acetate Buffer																
	cont	80	40	20	10	5	2.5	1.25	0.625	0.3125	0.1563	0.0781	0.0391	0.0195	0.0098	0.0049
MABL2(intact)	-	+	+	+	+	+	+	+	+	+	+	+	-	-	-	-

[0173]

[Effect of the invention]

The modified antibodies of the invention have an agonist action capable of transducing a signal into cells by crosslinking a cell surface molecule(s) and are advantageous in that the permeability to tissues and tumors is high due to the lowered molecular size compared with antibody molecule (whole IgG). The modified antibodies have remarkably higher activity compared with the original antibodies, which is attributable to that the modified antibodies are in a shape closer to a ligand compared with original antibodies. Therefore the modified antibodies can be used as signal-transducing agonists. The modification of antibody molecule results in the reduction of side effects caused by intercellular crosslinking and provides novel medicines inducing only required action by crosslinking a cell surface molecule(s). Medical preparations containing as active ingredient the modified antibody of the invention are useful as preventives and/or remedies for cancers, inflammation, hormone disorders and blood diseases, for example, leukemia, malignant lymphoma, aplastic anemia, myelodysplasia syndrome and polycythemia vera.

g G)と比較して低分子化が達成されているため、組織、腫瘍への移行性に優れているという特徴を有している。さらに本発明の改変抗体は、元のモノクローナル抗体と比較して顕著に高い活性を有しているが、これは本発明の改変抗体が抗体分子に比べてよりリガンドに近い形態であるためと考えられる。従って、当該改変抗体はシグナル伝達アゴニストとして使用することができ、そして抗体分子を本発明の改変抗体にすることにより、細胞間の架橋などによる副作用を軽減し、且つ細胞表面上の分子を架橋して所望の作用のみを誘起しうる新規な医薬品を提供される。本発明の改変抗体を有効成分とする医薬製剤は、癌、炎症、ホルモン異常、並びに白血病、悪性リンパ腫、再生不良性貧血、骨髓異形成症候群および真性多血症などの血液疾患の予防及び／又は治療薬として有用である。

【0174】

【配列表】

SEQUENCE LISTING

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<120> A polypeptide inducible apoptosis

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Ala Ser Ser Ser Asp Val Val Met Thr Gln Thr Pro Leu Ser Leu

20 25 30

cct gtc agt ctt gga gat caa gcc tcc atc tct tgc aga tct agt 135

Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser

35 40 45

cag agc ctt cta cac agt aaa gga aac acc tat tta caa tgg tac 180

Gln Ser Leu Leu His Ser Lys Gly Asn Thr Tyr Leu Gln Trp Tyr

50 55 60

cta cag aag cca ggc cag tct cca aag ctc ctg atc tac aaa gtt 225

Leu Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val

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Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly

80 85 90

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Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu

95 100 105

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Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr

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20

25

30

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35

40

45

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50

55

60

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整理番号=DOJ-5464

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95

100

105

gcc tct gag gac tct gcg gtc tac tac tgt gca aga ggg ggt tac 360

Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr

110

115

120

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Tyr Ser Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser

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tca g 409

Ser

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<212> DNA

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<220>

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15

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Gly Ser Ser Ser Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu

20

25

30

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Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser

35

40

45

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50

55

60

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Leu Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val

65

70

75

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80

85

90

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Ser Val Thr Asp Phe Thr Leu Met Ile Ser Arg Val Glu Ala Glu

95

100

105

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Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr

110

115

120

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Gly Val His Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu

20

25

30

gta aag cct ggg gct tca gtg aag atg tcc tgc aag gct tct gga 135

Val Lys Pro Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly

35

40

45

tac acc ttc gct aac cat gtt att cac tgg gtg aag cag aag cca 180

Tyr Thr Phe Ala Asn His Val Ile His Trp Val Lys Gln Lys Pro

50

55

60

ggg cag ggc ctt gag tgg att gga tat att tat cct tac aat gat 225

Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro Tyr Asn Asp

65

70

75

ggt act aag tat aat gag aag ttc aag gac aag gcc act ctg act 270

Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp Lys Ala Thr Leu Thr

80

85

90

tca gac aaa tcc tcc acc aca gcc tac atg gac ctc agc agc ctg 315

Ser Asp Lys Ser Ser Thr Thr Ala Tyr Met Asp Leu Ser Ser Leu

95

100

105

gcc tct gag gac tct gcg gtc tat tac tgt gca aga ggg ggt tac 360

Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr

110

115

120

tat act tac gac gac tgg ggc caa ggc acc act ctc aca gtc tcc 405

Tyr Thr Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser

125

130

135

tca g

409

Ser

<210> 9

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 9

cccaagcttc caccatgaag ttgcctgtta gg 32

<210> 10

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 10

cccaagcttc caccatggaa tggagctgga ta 32

<210> 11

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 11

cgcggatcca ctcacgtttt atttccagct tggt 34

<210> 12

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 12

cgcggatcca ctcacctgag gagactgtga gagt 34

<210> 13

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 13

catgccatgg cgcaggtcca gctgcagcag 30

<210> 14

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 14

accaccacct gaggagactg tgagagt 27

<210> 15

整理番号=D O J - 5 4 6 4

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 15

gtctcctcag gtggtggtgg ttcgggt 27

<210> 16

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 16

cacaacatcc gatccgccac cacccga 27

<210> 17

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 17

ggcggatcgg atgttgtgat gacccaa 27

<210> 18

<211> 57

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 18

ccggaattct cattatttat cgtcatcgtc ttgttagtct ttattttcca gcttggt 57

<210> 19

<211> 45

<212> DNA

<213> Artificial Sequence

<220>

<223> Linker amino acid sequence and nucleotide sequence

<400> 19

ggt ggt ggt ggt tcg ggt ggt ggt ggt tcg ggt ggt ggc gga tcg 45
Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
5 10 15

<210> 20

<211> 828

<212> DNA

<213> Mus

<220>

<221>CDS

<222>(1)...(826)

<223> pscM1. MABL1-scFv

<400> 20

atg aaa tac cta ttg cct acg gca gcc gct gga ttg tta tta ctc 45
Met Lys Tyr Leu Leu Pro Thr Ala Ala Ala Gly Leu Leu Leu Leu
5 10 15

gct gcc caa cca gcc atg gcg cag gtc cag ctg cag cag tct gga 90

Ala Ala Gln Pro Ala Met Ala Gln Val Gln Leu Gln Gln Ser Gly

20

25

30

cct gac ctg gta aag cct ggg gct tca gtg aag atg tcc tgc aag 135

Pro Asp Leu Val Lys Pro Gly Ala Ser Val Lys Met Ser Cys Lys

35

40

45

gct tct gga tac acc ttc gtt aac cat gtt atg cac tgg gtg aag 180

Ala Ser Gly Tyr Thr Phe Val Asn His Val Met His Trp Val Lys

50

55

60

cag aag cca ggg cag ggc ctt gag tgg att gga tat att tat cct 225

Gln Lys Pro Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro

65

70

75

tac aat gat ggt act aag tac aat gag aag ttc aag ggc aag gcc 270

Tyr Asn Asp Gly Thr Lys Tyr Asn Glu Lys Phe Lys Gly Lys Ala

80

85

90

aca ctg act tca gag aaa tcc tcc agc gca gcc tac atg gag ctc 315

Thr Leu Thr Ser Glu Lys Ser Ser Ser Ala Ala Tyr Met Glu Leu

95

100

105

agc agc ctg gcc tct gag gac tct gcg gtc tac tac tgt gca aga 360

Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg

110

115

120

ggg ggt tac tat agt tac gac gac tgg ggc caa ggc acc act ctc 405

Gly Gly Tyr Tyr Ser Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu

125

130

135

aca gtc tcc tca ggt ggt ggt ggt tcg ggt ggt ggt ggt tcg ggt 450

Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly

140

145

150

ggt ggc gga tcg gat gtt gtg atg acc caa act cca ctc tcc ctg 495

Gly Gly Gly Ser Asp Val Val Met Thr Gln Thr Pro Leu Ser Leu

155	160	165
cct gtc agt ctt gga gat caa gcc tcc atc tct tgc aga tct agt	540	
Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser		
170	175	180
cag agc ctt cta cac agt aaa gga aac acc tat tta caa tgg tac	585	
Gln Ser Leu Leu His Ser Lys Gly Asn Thr Tyr Leu Gln Trp Tyr		
185	190	195
cta cag aag cca ggc cag tct cca aag ctc ctg atc tac aaa gtt	630	
Leu Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val		
200	205	210
tcc aac cga ttt tct ggg gtc cca gac agg ttc agt ggc agt gga	675	
Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly		
215	220	225
tca ggg aca gat ttc aca ctc aag atc agc aga gtg gag gct gag	720	
Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu		
230	235	240
gat ctg gga gtt tat ttc tgc tct caa agt aca cat gtt ccg tac	765	
Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr		
245	250	255
acg tcc gga ggg ggg acc aag ctg gaa ata aaa gac tac aaa gac	810	
Thr Ser Gly Gly Gly Thr Lys Leu Glu Ile Lys Asp Tyr Lys Asp		
260	265	270
gat gac gat aaa taa tga		828
Asp Asp Asp Lys		

<210> 21

<211> 31

<212> DNA

<213> Artificial Sequence

整理番号=DOJ-5464

<220>

<223> PCR primer

<400> 21

acgcgtcgac tcccaggtcc agctgcagca g 31

<210> 22

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 22

gaaggtgtat ccagaagc 18

<210> 23

<211> 819

<212> DNA

<213> Mus

<220>

<221> CDS

<222>(1)...(813)

<223> pCHOM1. MABL1-scFv

<400> 23

atg gga tgg agc tgt atc atc ctc ttc ttg gta gca aca gct aca 45

Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr

5

10

15

ggt gtc gac tcc cag gtc cag ctg cag cag tct gga cct gac ctg 90

Gly Val Asp Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Asp Leu

20

25

30

gta aag cct ggg gct tca gtg aag atg tcc tgc aag gct tct gga	135
Val Lys Pro Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly	
35	40 45
tac acc ttc gtt aac cat gtt atg cac tgg gtg aag cag aag cca	180
Tyr Thr Phe Val Asn His Val Met His Trp Val Lys Gln Lys Pro	
50	55 60
ggg cag ggc ctt gag tgg att gga tat att tat cct tac aat gat	225
Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro Tyr Asn Asp	
65	70 75
ggt act aag tac aat gag aag ttc aag ggc aag gcc aca ctg act	270
Gly Thr Lys Tyr Asn Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr	
80	85 90
tca gag aaa tcc tcc agc gca gcc tac atg gag ctc agc agc ctg	315
Ser Glu Lys Ser Ser Ser Ala Ala Tyr Met Glu Leu Ser Ser Leu	
95	100 105
gcc tct gag gac tct gcg gtc tac tac tgt gca aga ggg ggt tac	360
Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr	
110	115 120
tat agt tac gac gac tgg ggc caa ggc acc act ctc aca gtc tcc	405
Tyr Ser Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser	
125	130 135
tca ggt ggt ggt ggt tcg ggt ggt ggt ggt tcg ggt ggt ggc gga	450
Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly	
140	145 150
tcg gat gtt gtg atg acc caa act cca ctc tcc ctg cct gtc agt	495
Ser Asp Val Val Met Thr Gln Thr Pro Leu Ser Leu Pro Val Ser	
155	160 165
ctt gga gat caa gcc tcc atc tct tgc aga tct agt cag agc ctt	540
Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu	

170	175	180	
cta cac agt aaa gga aac acc tat tta caa tgg tac cta cag aag	585		
Leu His Ser Lys Gly Asn Thr Tyr Leu Gln Trp Tyr Leu Gln Lys			
185	190	195	
cca ggc cag tct cca aag ctc ctg atc tac aaa gtt tcc aac cga	630		
Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg			
200	205	210	
TTT TCT GGG GTC CCA GAC AGG TTC AGT GGC AGT GGA TCA GGG ACA	675		
Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr			
215	220	225	
gat ttc aca ctc aag atc agc aga gtg gag gct gag gat ctg gga	720		
Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly			
230	235	240	
gtt tat ttc tgc tct caa agt aca cat gtt ccg tac acg tcc gga	765		
Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr Thr Ser Gly			
245	250	255	
ggg ggg acc aag ctg gaa ata aaa gac tac aaa gac gat gac gat	810		
Gly Gly Thr Lys Leu Glu Ile Lys Asp Tyr Lys Asp Asp Asp Asp			
260	265	270	
aaa taa tga			819
Lys			

<210> 24

<211> 828

<212> DNA

<213> Mus

<220>

<221> CDS

<222> (1)...(822)

<223> pscM2. MABL2-scFv

<400> 24

atg aaa tac cta ttg cct acg gca gcc gct gga ttg tta tta ctc 45

Met Lys Tyr Leu Leu Pro Thr Ala Ala Ala Gly Leu Leu Leu Leu

5 10 15

gct gcc caa cca gcc atg gcg cag gtc cag ctg cag cag tct gga 90

Ala Ala Gln Pro Ala Met Ala Gln Val Gln Leu Gln Gln Ser Gly

20 25 30

cct gaa ctg gta aag cct ggg gct tca gtg aag atg tcc tgc aag 135

Pro Glu Leu Val Lys Pro Gly Ala Ser Val Lys Met Ser Cys Lys

35 40 45

gct tct gga tac acc ttc gct aac cat gtt att cac tgg gtg aag 180

Ala Ser Gly Tyr Thr Phe Ala Asn His Val Ile His Trp Val Lys

50 55 60

cag aag cca ggg cag ggc ctt gag tgg att gga tat att tat cct 225

Gln Lys Pro Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro

65 70 75

tac aat gat ggt act aag tat aat gag aag ttc aag gac aag gcc 270

Tyr Asn Asp Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp Lys Ala

80 85 90

act ctg act tca gac aaa tcc tcc acc aca gcc tac atg gac ctc 315

Thr Leu Thr Ser Asp Lys Ser Ser Thr Thr Ala Tyr Met Asp Leu

95 100 105

agc agc ctg gcc tct gag gac tct gcg gtc tat tac tgt gca aga 360

Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg

110 115 120

ggg ggt tac tat act tac gac gac tgg ggc caa ggc acc act ctc 405

Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu

125 130 135

aca gtc tcc tca ggt ggt ggt ggt tgc ggt ggt ggt ggt tgc ggt 450
 Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
 140 145 150
 ggt ggc gga tgc gat gtt gtg atg acc caa agt cca ctc tcc ctg 495
 Gly Gly Gly Ser Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu
 155 160 165
 cct gtc agt ctt gga gat caa gcc tcc atc tct tgc aga tca agt 540
 Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser
 170 175 180
 cag agc ctt gtg cac agt aat gga aag acc tat tta cat tgg tac 585
 Gln Ser Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr
 185 190 195
 ctg cag aag cca ggc cag tct cca aaa ctc ctg atc tac aaa gtt 630
 Leu Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val
 200 205 210
 tcc aac cga ttt tct ggg gtc cca gac agg ttc agt ggc agt gga 675
 Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly
 215 220 225
 tca gtg aca gat ttc aca ctc atg atc agc aga gtg gag gct gag 720
 Ser Val Thr Asp Phe Thr Leu Met Ile Ser Arg Val Glu Ala Glu
 230 235 240
 gat ctg gga gtt tat ttc tgc tct caa agt aca cat gtt ccg tac 765
 Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr
 245 250 255
 acg ttc gga ggg ggg acc aag ctg gaa ata aaa gac tac aaa gac 810
 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Asp Tyr Lys Asp
 260 265 270
 gat gac gat aaa taa tga 828
 Asp Asp Asp Lys

<210> 25

<211> 819

<212> DNA

<213> Mus

<220>

<221> CDS

<222> (1)...(813)

<223> pCHOM2. MABL2-scFv

<400> 25

atg gga tgg agc tgt atc atc ctc ttc ttg gta gca aca gct aca 45

Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr

5

10

15

ggt gtc gac tcc cag gtc cag ctg cag cag tct gga cct gaa ctg 90

Gly Val Asp Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu

20

25

30

gta aag cct ggg gct tca gtg aag atg tcc tgc aag gct tct gga 135

Val Lys Pro Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly

35

40

45

tac acc ttc gct aac cat gtt att cac tgg gtg aag cag aag cca 180

Tyr Thr Phe Ala Asn His Val Ile His Trp Val Lys Gln Lys Pro

50

55

60

ggg cag ggc ctt gag tgg att gga tat att tat cct tac aat gat 225

Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro Tyr Asn Asp

65

70

75

ggt act aag tat aat gag aag ttc aag gac aag gcc act ctg act 270

Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp Lys Ala Thr Leu Thr

80

85

90

tca gac aaa tcc tcc acc aca gcc tac atg gac ctc agc agc ctg 315

Ser Asp Lys Ser Ser Thr Thr Ala Tyr Met Asp Leu Ser Ser Leu
 95 100 105
 gcc tct gag gac tct gcg gtc tat tac tgt gca aga ggg ggt tac 360
 Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr
 110 115 120
 tat act tac gac gac tgg ggc caa ggc acc act ctc aca gtc tcc 405
 Tyr Thr Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser
 125 130 135
 tca ggt ggt ggt ggt tcg ggt ggt ggt ggt tcg ggt ggt ggc gga 450
 Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly
 140 145 150
 tcg gat gtt gtg atg acc caa agt cca ctc tcc ctg cct gtc agt 495
 Ser Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro Val Ser
 155 160 165
 ctt gga gat caa gcc tcc atc tct tgc aga tca agt cag agc ctt 540
 Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu
 170 175 180
 gtg cac agt aat gga aag acc tat tta cat tgg tac ctg cag aag 585
 Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys
 185 190 195
 cca ggc cag tct cca aaa ctc ctg atc tac aaa gtt tcc aac cga 630
 Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg
 200 205 210
 ttt tct ggg gtc cca gac agg ttc agt ggc agt gga tca gtg aca 675
 Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Val Thr
 215 220 225
 gat ttc aca ctc atg atc agc aga gtg gag gct gag gat ctg gga 720
 Asp Phe Thr Leu Met Ile Ser Arg Val Glu Ala Glu Asp Leu Gly
 230 235 240

gtt tat ttc tgc tct caa agt aca cat gtt ccg tac acg ttc gga 765

Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly

245

250

255

ggg ggg acc aag ctg gaa ata aaa gac tac aaa gac gat gac gat 810

Gly Gly Thr Lys Leu Glu Ile Lys Asp Tyr Lys Asp Asp Asp Asp

260

265

270

aaa taa tga

819

Lys

<210> 26

<211> 456

<212> DNA

<213> Mus

<220>

<221> CDS

<222> (1)...(450)

<223> pCHO-shIAP. Soluble human IAP

<400> 26

atg tgg ccc ctg gta gcg gcg ctg ttg ctg ggc tcg gcg tgc tgc 45

Met Trp Pro Leu Val Ala Ala Leu Leu Leu Gly Ser Ala Cys Cys

5

10

15

gga tca gct cag cta cta ttt aat aaa aca aaa tct gta gaa ttc 90

Gly Ser Ala Gln Leu Leu Phe Asn Lys Thr Lys Ser Val Glu Phe

20

25

30

acg ttt tgt aat gac act gtc gtc att cca tgc ttt gtt act aat 135

Thr Phe Cys Asn Asp Thr Val Val Ile Pro Cys Phe Val Thr Asn

35

40

45

atg gag gca caa aac act act gaa gta tac gta aag tgg aaa ttt 180

Met Glu Ala Gln Asn Thr Thr Glu Val Tyr Val Lys Trp Lys Phe

50	55	60	
aaa gga aga gat att tac acc ttt gat gga gct cta aac aag tcc	225		
Lys Gly Arg Asp Ile Tyr Thr Phe Asp Gly Ala Leu Asn Lys Ser			
65	70	75	
act gtc ccc act gac ttt agt agt gca aaa att gaa gtc tca caa	270		
Thr Val Pro Thr Asp Phe Ser Ser Ala Lys Ile Glu Val Ser Gln			
80	85	90	
tta cta aaa gga gat gcc tct ttg aag atg gat aag agt gat gct	315		
Leu Leu Lys Gly Asp Ala Ser Leu Lys Met Asp Lys Ser Asp Ala			
95	100	105	
gtc tca cac aca gga aac tac act tgt gaa gta aca gaa tta acc	360		
Val Ser His Thr Gly Asn Tyr Thr Cys Glu Val Thr Glu Leu Thr			
110	115	120	
aga gaa ggt gaa acg atc atc gag cta aaa tat cgt gtt gtt tca	405		
Arg Glu Gly Glu Thr Ile Ile Glu Leu Lys Tyr Arg Val Val Ser			
125	130	135	
tgg ttt tct cca aat gaa aat gac tac aag gac gac gat gac aag	450		
Trp Phe Ser Pro Asn Glu Asn Asp Tyr Lys Asp Asp Asp Asp Lys			
140	145	150	
tga tag			456

<210> 27

<211> 46

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 27

ggaattccat atgcaagtgc aacttcaaca gtctggacct gaactg 46

<210> 28

<211> 31

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 28

ggaattctca ttattttatt tccagcttgg t 31

<210> 29

<211> 741

<212> DNA

<213> Mus

<220>

<221> CDS

<222> (1)...(735)

<223> pscM2DEm02. MABL2-scFv

<400> 29

atg caa gtg caa ctt caa cag tct gga cct gaa ctg gta aag cct 45

Met Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro

5

10

15

ggg gct tca gtg aag atg tcc tgc aag gct tct gga tac acc ttc 90

Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe

20

25

30

gct aac cat gtt att cac tgg gtg aag cag aag cca ggg cag ggc 135

Ala Asn His Val Ile His Trp Val Lys Gln Lys Pro Gly Gln Gly

35

40

45

ctt gag tgg att gga tat att tat cct tac aat gat ggt act aag 180

Leu Glu Trp Ile Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys
 50 55 60
 tat aat gag aag ttc aag gac aag gcc act ctg act tca gac aaa 225
 Tyr Asn Glu Lys Phe Lys Asp Lys Ala Thr Leu Thr Ser Asp Lys
 65 70 75
 tcc tcc acc aca gcc tac atg gac ctc agc agc ctg gcc tct gag 270
 Ser Ser Thr Thr Ala Tyr Met Asp Leu Ser Ser Leu Ala Ser Glu
 80 85 90
 gac tct gcg gtc tat tac tgt gca aga ggg ggt tac tat act tac 315
 Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr
 95 100 105
 gac gac tgg ggc caa ggc acc act ctc aca gtc tcc tca ggt ggt 360
 Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser Ser Gly Gly
 110 115 120
 ggt ggt tcg ggt ggt ggt ggt tcg ggt ggt ggc gga tcg gat gtt 405
 Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Asp Val
 125 130 135
 gtg atg acc caa agt cca ctc tcc ctg cct gtc agt ctt gga gat 450
 Val Met Thr Gln Ser Pro Leu Ser Leu Pro Val Ser Leu Gly Asp
 140 145 150
 caa gcc tcc atc tct tgc aga tca agt cag agc ctt gtg cac agt 495
 Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser
 155 160 165
 aat gga aag acc tat tta cat tgg tac ctg cag aag cca ggc cag 540
 Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln
 170 175 180
 tct cca aaa ctc ctg atc tac aaa gtt tcc aac cga ttt tct ggg 585
 Ser Pro Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly
 185 190 195

gtc cca gac agg ttc agt ggc agt gga tca gtg aca gat ttc aca 630

Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Val Thr Asp Phe Thr

200

205

210

ctc atg atc agc aga gtg gag gct gag gat ctg gga gtt tat ttc 675

Leu Met Ile Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe

215

220

225

tgc tct caa agt aca cat gtt ccg tac acg ttc gga ggg ggg acc 720

Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gly Gly Thr

230

235

240

aag ctg gaa ata aaa taa tga

741

Lys Leu Glu Ile Lys

<210> 30

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 30

cagacagtgg ttcaaagt 18

<210> 31

<211> 72

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 31

cgcgctcgacc gatccgccc caccggaacc accaccaccc gaaccaccac caccttttat ttccag

cttg gt 72

<210> 32

<211> 1605

<212> DNA

<213> Mus

<220>

<221> CDS

<222> (1)...(1599)

<223> pCHOM2(Fv)2. MABL2-sc(Fv)2

<400> 32

atg gga tgg agc tgt atc atc ctc ttc ttg gta gca aca gct aca 45

Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr

5

10

15

ggt gtc gac tcc cag gtc cag ctg cag cag tct gga cct gaa ctg 90

Gly Val Asp Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu

20

25

30

gta aag cct ggg gct tca gtg aag atg tcc tgc aag gct tct gga 135

Val Lys Pro Gly Ala Ser Val Lys Met Ser Cys Lys Ala Ser Gly

35

40

45

tac acc ttc gct aac cat gtt att cac tgg gtg aag cag aag cca 180

Tyr Thr Phe Ala Asn His Val Ile His Trp Val Lys Gln Lys Pro

50

55

60

ggg cag ggc ctt gag tgg att gga tat att tat cct tac aat gat 225

Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro Tyr Asn Asp

65

70

75

ggt act aag tat aat gag aag ttc aag gac aag gcc act ctg act 270

Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp Lys Ala Thr Leu Thr

80

85

90

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tca gac aaa tcc tcc acc aca gcc tac atg gac ctc agc agc ctg	315
Ser Asp Lys Ser Ser Thr Thr Ala Tyr Met Asp Leu Ser Ser Leu	
95	100 105
gcc tct gag gac tct gcg gtc tat tac tgt gca aga ggg ggt tac	360
Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr	
110	115 120
tat act tac gac gac tgg ggc caa ggc acc act ctc aca gtc tcc	405
Tyr Thr Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser	
125	130 135
tca ggt ggt ggt ggt tcg ggt ggt ggt ggt tcg ggt ggt ggc gga	450
Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly	
140	145 150
tcg gat gtt gtg atg acc caa agt cca ctc tcc ctg cct gtc agt	495
Ser Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro Val Ser	
155	160 165
ctt gga gat caa gcc tcc atc tct tgc aga tca agt cag agc ctt	540
Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu	
170	175 180
gtg cac agt aat gga aag acc tat tta cat tgg tac ctg cag aag	585
Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys	
185	190 195
cca ggc cag tct cca aaa ctc ctg atc tac aaa gtt tcc aac cga	630
Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg	
200	205 210
ttt tct ggg gtc cca gac agg ttc agt ggc agt gga tca gtg aca	675
Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Val Thr	
215	220 225
gat ttc aca ctc atg atc agc aga gtg gag gct gag gat ctg gga	720
Asp Phe Thr Leu Met Ile Ser Arg Val Glu Ala Glu Asp Leu Gly	

230	235	240
gtt tat ttc tgc tct caa agt aca cat gtt ccg tac acg ttc gga	765	
Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly		
245	250	255
ggg ggg acc aag ctg gaa ata aaa ggt ggt ggt ggt tcg ggt ggt	810	
Gly Gly Thr Lys Leu Glu Ile Lys Gly Gly Gly Gly Ser Gly Gly		
260	265	270
ggt ggt tcg ggt ggt ggc gga tcg gtc gac tcc cag gtc cag ctg	855	
Gly Gly Ser Gly Gly Gly Gly Ser Val Asp Ser Gln Val Gln Leu		
275	280	285
cag cag tct gga cct gaa ctg gta aag cct ggg gct tca gtg aag	900	
Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Ala Ser Val Lys		
290	295	300
atg tcc tgc aag gct tct gga tac acc ttc gct aac cat gtt att	945	
Met Ser Cys Lys Ala Ser Gly Tyr Thr Phe Ala Asn His Val Ile		
305	310	315
cac tgg gtg aag cag aag cca ggg cag ggc ctt gag tgg att gga	990	
His Trp Val Lys Gln Lys Pro Gly Gln Gly Leu Glu Trp Ile Gly		
320	325	330
tat att tat cct tac aat gat ggt act aag tat aat gag aag ttc	1035	
Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn Glu Lys Phe		
335	340	345
aag gac aag gcc act ctg act tca gac aaa tcc tcc acc aca gcc	1080	
Lys Asp Lys Ala Thr Leu Thr Ser Asp Lys Ser Ser Thr Thr Ala		
350	355	360
tac atg gac ctc agc agc ctg gcc tct gag gac tct gcg gtc tat	1125	
Tyr Met Asp Leu Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr		
365	370	375
tac tgt gca aga ggg ggt tac tat act tac gac gac tgg ggc caa	1170	

Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln			
380	385		390
ggc acc act ctc aca gtc tcc tca ggt ggt ggt ggt tcg ggt ggt	1215		
Gly Thr Thr Leu Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly			
395	400		405
ggt ggt tcg ggt ggt ggc gga tcg gat gtt gtg atg acc caa agt	1260		
Gly Gly Ser Gly Gly Gly Gly Ser Asp Val Val Met Thr Gln Ser			
410	415		420
cca ctc tcc ctg cct gtc agt ctt gga gat caa gcc tcc atc tct	1305		
Pro Leu Ser Leu Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser			
425	430		435
tgc aga tca agt cag agc ctt gtg cac agt aat gga aag acc tat	1350		
Cys Arg Ser Ser Gln Ser Leu Val His Ser Asn Gly Lys Thr Tyr			
440	445		450
tta cat tgg tac ctg cag aag cca ggc cag tct cca aaa ctc ctg	1395		
Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu			
455	460		465
atc tac aaa gtt tcc aac cga ttt tct ggg gtc cca gac agg ttc	1440		
Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe			
470	475		480
agt ggc agt gga tca gtg aca gat ttc aca ctc atg atc agc aga	1485		
Ser Gly Ser Gly Ser Val Thr Asp Phe Thr Leu Met Ile Ser Arg			
485	490		495
gtg gag gct gag gat ctg gga gtt tat ttc tgc tct caa agt aca	1530		
Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr			
500	505		510
cat gtt ccg tac acg ttc gga ggg ggg acc aag ctg gaa ata aaa	1575		
His Val Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys			
515	520		525

gac tac aaa gac gat gac gat aaa taa tga

1605

Asp Tyr Lys Asp Asp Asp Asp Lys

530

<210> 33

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cacgacgtca ctcgagactg tgagagtggg gccttggccc 40

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<213> Artificial Sequence

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<223> PCR primer

<400> 37

cgcgtaatac gactcactat ag 22

<210> 38

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gcaattggac ctgttttate tcgagcttgg tccccctcc gaacgt 46

整理番号=DOJ-5464

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<400> 41

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整理番号=DOJ-5464

<212> DNA

<213> Artificial Sequence

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<211> 49

<212> DNA

<213> Artificial Sequence

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<223> PCR primer

<400> 45

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<210> 46

<211> 52

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 46

cagtctcgag tggtaggtggt ggtggtggtt ccgacgtcgt gatgacccaa ag 52

<210> 47

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 47

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<210> 48

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<220>

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MET Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly Val

gac tcc cag gtc cag ctg cag cag tct gga cct gaa ctg gta aag cct ggg 102
Asp Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly

gct tca gtg aag atg tcc tgc aag gct tct gga tac acc ttc gct aac cat 153
Ala Ser Val Lys MET Ser Cys Lys Ala Ser Gly Tyr Thr Phe Ala Asn His

gtt att cac tgg gtg aag cag aag cca ggg cag ggc ctt gag tgg att gga 204
Val Ile His Trp Val Lys Gln Lys Pro Gly Gln Gly Leu Glu Trp Ile Gly

tat att tat cct tac aat gat ggt act aag tat aat gag aag ttc aag gac 255
Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp

aag gcc act ctg act tca gac aaa tcc tcc acc aca gcc tac atg gac ctc 306
Lys Ala Thr Leu Thr Ser Asp Lys Ser Ser Thr Thr Ala Tyr MET Asp Leu

agc agc ctg gcc tct gag gac tct gcg gtc tat tac tgt gca aga ggg ggt 357
Ser Ser Leu Ala Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly

tac tat act tac gac gac tgg ggc caa ggc acc act ctc aca gtc tcg agt 408
Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln Gly Thr Thr Leu Thr Val Ser Ser

gac gtc gtg atg acc caa agt cca ctc tcc ctg cct gtc agt ctt gga gat 459
Asp Val Val MET Thr Gln Ser Pro Leu Ser Leu Pro Val Ser Leu Gly Asp

caa gcc tcc atc tct tgc aga tca agt cag agc ctt gtg cac agt aat gga 510

Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser Asn Gly

aag acc tat tta cat tgg tac ctg cag aag cca ggc cag tct cca aaa ctc 561

Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Lys Leu

ctg atc tac aaa gtt tcc aac cga ttt tct ggg gtc cca gac agg ttc agt 612

Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser

ggc agt gga tca gtg aca gat ttc aca ctc atg atc agc aga gtg gag gct 663

Gly Ser Gly Ser Val Thr Asp Phe Thr Leu MET Ile Ser Arg Val Glu Ala

gag gat ctg gga gtt tat ttc tgc tct caa agt aca cat gtt ccg tac acg 714

Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Tyr Thr

ttc gga ggg ggg acc aag ctg gaa ata aaa gac tac aaa gac gat gac gat 765

Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Asp Tyr Lys Asp Asp Asp Asp

aaa taa tga gga tcc

780

Lys

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<210> 52

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caagctcgag ataaaatccg gaggtggtgg tggccaggtc caattgcagc agtc 54

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<210> 53

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<213> Artificial Sequence

<220>

<223> PCR primer

<400> 53

caagctcgag ataaaatccg gaggtggtgg tggtagccag gtccaattgc agcagtc 57

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<211> 780

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<213> Mus

<220>

<221> CDS

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<223> CF2LH-0/pCOS1. MABL2-scFv<LH-0>

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MET Lys Leu Pro Val Arg Leu Leu Val Leu MET Phe Trp Ile Pro Gly Ser

agc agt gat gtt gtg atg acc caa agt cca ctc tcc ctg cct gtc agt ctt 102

Ser Ser Asp Val Val MET Thr Gln Ser Pro Leu Ser Leu Pro Val Ser Leu

gga gat caa gcc tcc atc tct tgc aga tca agt cag agc ctt gtg cac agt 153

Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser

aat gga aag acc tat tta cat tgg tac ctg cag aag cca ggc cag tct cca 204

Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro

aaa ctc ctg atc tac aaa gtt tcc aac cga ttt tct ggg gtc cca gac agg 255
Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg

ttc agt ggc agt gga tca gtg aca gat ttc aca ctc atg atc agc aga gtg 306
Phe Ser Gly Ser Gly Ser Val Thr Asp Phe Thr Leu MET Ile Ser Arg Val

gag gct gag gat ctg gga gtt tat ttc tgc tct caa agt aca cat gtt ccg 357
Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro

tac acg ttc gga ggg ggg acc aag ctc gag ata aaa cag gtc caa ttg cag 408
Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Gln Val Gln Leu Gln

cag tct gga cct gaa ctg gta aag cct ggg gct tca gtg aag atg tcc tgc 459
Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Ala Ser Val Lys MET Ser Cys

aag gct tct gga tac acc ttc gct aac cat gtt att cac tgg gtg aag cag 510
Lys Ala Ser Gly Tyr Thr Phe Ala Asn His Val Ile His Trp Val Lys Gln

aag cca ggg cag ggc ctt gag tgg att gga tat att tat cct tac aat gat 561
Lys Pro Gly Gln Gly Leu Glu Trp Ile Gly Tyr Ile Tyr Pro Tyr Asn Asp

ggc act aag tat aat gag aag ttc aag gac aag gcc act ctg act tca gac 612
Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp Lys Ala Thr Leu Thr Ser Asp

aaa tcc tcc acc aca gcc tac atg gac ctc agc agc ctg gcc tct gag gac 663
Lys Ser Ser Thr Thr Ala Tyr MET Asp Leu Ser Ser Leu Ala Ser Glu Asp

tct gcg gtc tat tac tgt gca aga ggg ggt tac tat act tac gac gac tgg 714

Ser Ala Val Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp

ggc caa ggc acc act etc aca gtc tcc tca gac tac aaa gac gat gac gat 765

Gly Gln Gly Thr Thr Leu Thr Val Ser Ser Asp Tyr Lys Asp Asp Asp Asp

aaa taa tga gga tcc

780

Lys

【図面の簡単な説明】

【図1】

ヒトIgG1抗体が、ヒトIAPを発現するL1210細胞(hIAP/L1210)に結合しないことを示すフローサイトメトリーの結果を示す図である。

【図2】

キメラMABL-1抗体が、ヒトIAPを発現するL1210細胞(hIAP/L1210)に特異的に結合することを示すフローサイトメトリーの結果を示す図である。

【図3】

キメラMABL-2抗体が、ヒトIAPを発現するL1210細胞(hIAP/L1210)に特異的に結合することを示すフローサイトメトリーの結果を示す図である。

【図4】

本発明にかかる一本鎖Fvの作成方法を模式的に示す図である。

【図5】

本発明の一本鎖FvをコードするDNAを、大腸菌にて発現させるために使用可能な発現プラスミドの一例の構造を示す。

【図6】

本発明の一本鎖FvをコードするDNAを、哺乳動物細胞にて発現させるために使用する発現プラスミドの一例の構造を示す。

[EXPLANATION OF DRAWINGS]

Figure 1 shows the result of flow cytometry, illustrating that human IgG antibody does not bind to L1210 cells expressing human IAP (hIAP/L1210).

5 Figure 2 shows the result of flow cytometry, illustrating that the chimera MABL-1 antibody specifically binds to L1210 cells expressing human IAP (hIAP/L1210).

10 Figure 3 shows the result of flow cytometry, illustrating that the chimera MABL-2 antibody specifically binds to L1210 cells expressing human IAP (hIAP/L1210).

Figure 4 schematically illustrates the process for producing the single chain Fv according to the present invention.

15 Figure 5 illustrates a structure of an expression plasmid which can be used to express a DNA encoding the single chain Fv of the invention in E. coli.

Figure 6 illustrates a structure of an expression plasmid which is used to express a DNA encoding the single chain Fv of the invention in mammalian cells.

20 Figure 7 shows a photograph showing the result of western blotting in Example 5.4. From the left, a molecular weight marker (which indicates 97.4, 66, 45, 31, 21.5 and 14.5 kDa from the top), the culture supernatant of pCHO1-introduced COS7 cells and the culture supernatant of pCHOM2-introduced COS7 cells. It illustrates that the reconstructed
25 single chain Fv of the antibody MABL-2 (arrow) is contained in the culture supernatant of the pCHOM2-introduced cells.

Figure 8 shows the result of flow cytometry, illustrating that an antibody in the culture supernatant of pCHO1/COS7 cell as a control does not bind to pCOS1/L1210 cell as a control.

5 Figure 9 shows the result of flow cytometry, illustrating that an antibody in the culture supernatant of MABL2-scFv/COS7 cells does not bind to pCOS1/L1210 cells as a control.

10 Figure 10 shows the result of flow cytometry, illustrating that an antibody in the culture supernatant of pCOS1/COS7 cells as a control does not bind to hIAP/L1210 cells.

15 Figure 11 shows the result of flow cytometry, illustrating that an antibody in the culture supernatant of MABL2-scFv/COS7 cells specifically binds to hIAP/L1210 cells.

20 Figure 12 shows the result of the competitive ELISA in Example 5.6, wherein the binding activity of the single chain Fv of the invention (MABL2-scFv) to the antigen is demonstrated in terms of the inhibition of binding of the mouse monoclonal antibody MABL-2 to the antigen as an index, in comparison with the culture supernatant of pCHO1/COS7 cells as a control.

25 Figure 13 shows the results of the apoptosis-inducing effect in Example 5.7, illustrating that the antibody in the culture supernatant of pCHO1/COS7 cells as a

control does not induce the apoptosis of pCOS1/L1210 cells as a control.

Figure 14 shows the results of the apoptosis-inducing effect in Example 5.7, illustrating that the antibody in the culture supernatant of MABL2-scFv/COS7 cells does not induce apoptosis of pCOS1/L1210 cells as a control.

Figure 15 shows the results of the apoptosis-inducing effect in Example 5.7, illustrating that the antibody in the culture supernatant of pCHO1/COS7 cells as a control does not induce apoptosis of hIAP/L1210 cells.

Figure 16 shows the results of the apoptosis-inducing effect in Example 5.7, illustrating that the antibody in the culture supernatant of MABL2-scFv/COS7 cells specifically induces apoptosis of hIAP/L1210 cells.

Figure 17 shows the results of the apoptosis-inducing effect in Example 5.7, illustrating that the antibody in the culture supernatant of pCHO1/COS7 cells as a control does not induce apoptosis of CCRF-CEM cells (at 50% of the final concentration).

Figure 18 shows the results of the apoptosis-inducing effect in Example 5.7, illustrating that the antibody in the culture supernatant of MABL2-scFv/COS7 cells specifically induces apoptosis of CCRF-CEM cells (at 50% of the final concentration).

Figure 19 shows the chromatogram obtained in the purification of the single chain Fv derived from the antibody MABL-2 produced by the CHO cells in Example 5.9,

illustrating that fraction A and fraction B were obtained as the major peaks when the fraction from Blue-sepharose column was purified with hydroxyapatite column.

Figure 20 shows the results of purification by gel filtration of fraction A and fraction B obtained in Example 5.9-(2), illustrating that the major peaks (AI and BI, respectively) were eluted from fraction A at approximately 36 kD of the apparent molecular weight and from fraction B at approximately 76 kD.

Figure 21 is the analysis on SDS-PAGE of the fractions obtained in the purification of the single chain Fv derived from the antibody MABL-2 produced by the CHO cells in Example 5.9, illustrating that a single band of approximately 35 kD of molecular weight was observed in both fractions.

Figure 22 shows the results of analysis of fractions AI and BI obtained by gel filtration in the purification of the single chain Fv derived from the antibody MABL-2 produced by the CHO cells, wherein fraction AI comprises monomer and fraction BI comprises dimer.

Figure 23 illustrates a structure of an expression plasmid which can be used to express a DNA encoding the single chain Fv of the invention in E. coli.

Figure 24 shows the results of purification on the gel filtration column of crude products of the single chain Fv polypeptide derived from the antibody MABL-2 produced by E. coli obtained in Example 5.12, wherein each peak

indicates monomer or dimer, respectively, of the single chain Fv produced by E. coli.

Figure 25 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that mouse IgG antibody as a control does not induce apoptosis of hIAP/L1210 cells (the final concentration of 3 $\mu\text{g/ml}$).

Figure 26 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that the dimer of MABL2-scFv produced by the CHO cells remarkably induces apoptosis of hIAP/L1210 cells (the final concentration of 3 $\mu\text{g/ml}$).

Figure 27 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that the dimer of MABL2-scFv produced by E. coli remarkably induces apoptosis of hIAP/L1210 cells (the final concentration of 3 $\mu\text{g/ml}$).

Figure 28 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that apoptosis induction to hIAP/L1210 cells by the MABL2-scFv monomer produced by the CHO cells is the same level as that of the control (the final concentration of 3 $\mu\text{g/ml}$).

Figure 29 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that apoptosis induction to hIAP/L1210 cells of the MABL2-scFv monomer produced by E. coli is the same level as that of control (the final concentration of 3 $\mu\text{g/ml}$).

Figure 30 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that mouse IgG antibody used as a control does not induce apoptosis of hIAP/L1210 cells even when anti-FLAG antibody is added (the final concentration of 3 μ g/ml).

Figure 31 shows the results of the apoptosis-inducing effect in Example 5.13, illustrating that MABL2-scFv monomer produced by the CHO cells remarkably induces apoptosis of hIAP/L1210 cells when anti-FLAG antibody is added (the final concentration of 3 μ g/ml).

Figure 32 shows the results of quantitative measurement of human IgG in the serum of a human myeloma cell line KPMM2-transplanted mouse, indicating amounts of human IgG produced by the human myeloma cells in the mouse. It illustrates that the dimer of scFv/CHO remarkably inhibited growth of the KPMM2 cells.

Figure 33 shows the survival time of the mouse after the transplantation of tumor, illustrating that the scFv/CHO dimer-administered group elongated remarkably the survival time.

Figure 34 illustrates a structure of an expression plasmid which expresses a reconstructed polypeptide_
[sc(Fv)₂] comprising two H chain V regions and two L chain V regions derived from the antibody MABL-2.

Figure 35 illustrates a structure of a plasmid which expresses a scFv (HL type) wherein the V regions are

linked in the manner of [H chain]-[L chain] without a peptide linker.

Figure 36 illustrates a structure of the HL-type polypeptide and amino acid sequences of peptide linkers.

5 Figure 37 illustrates a structure of a plasmid which expresses a scFv (LH type) wherein the V regions are linked in the manner of [L chain]-[H chain] without a peptide linker.

10 Figure 38 illustrates a structure of the LH-type polypeptide and amino acid sequences of peptide linkers.

15 Figure 39 shows the results of the western blotting in Example 6.4, illustrating that the reconstructed polypeptide sc(Fv)₂ comprising two H chain V regions and two L chain V regions, and the MABL2-scFv having peptide linkers with different length are expressed.

20 Figure 40 shows the results of flow cytometry using the culture supernatant of COS7 cells prepared in Example 6.3 (1), illustrating that the MABL2-scFv and sc(Fv)₂ having peptide linkers with different length have high affinities against human IAP.

 Figure 41 shows the results of the apoptosis-inducing effect in Example 6.6, illustrating that the scFv <HL3, 4, 6, 7, LH3, 4, 6 and 7> and the sc(Fv)₂ remarkably induce cell death of hIAP/L1210 cells.

25 Figure 42 shows the results of the evaluation of antigen binding capacity in Example 6.10, illustrating that

the dimer of scFv <HL5> and sc(Fv)₂ have high affinities against human IAP.

Figure 43 shows the results of the in vitro apoptosis-inducing effect in Example 6.11, illustrating that the dimer of scFv <HL5> and the sc(Fv)₂ induce apoptosis of hIAP/L1210 cells and CCRF-CEM cells in concentration-dependent manner.

[Document] Abstract

[Summary]

To provide a modified antibody comprising two or more H chain V regions and two or more L chain V regions of a monoclonal antibody and can act as an agonist transducing a signal into cells by crosslinking a cell surface molecule(s).

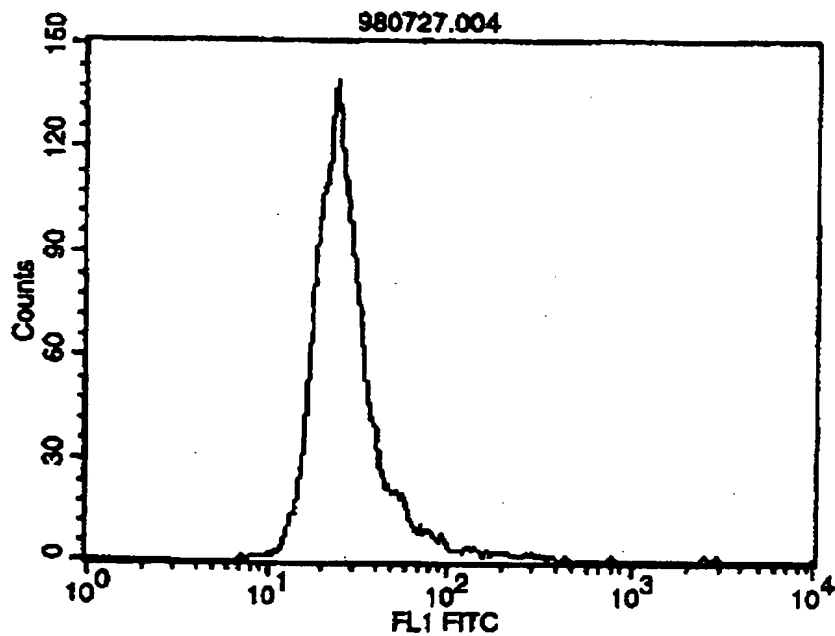
[Means to solve the problem]

The modified antibodies contain two or more H chain V regions and two or more L chain V regions of a monoclonal antibody and which combine with a cell surface molecule(s) and transduce a signal into cells, thereby can serve as an agonist. The modified antibodies can be used as signal-transducing agonists. Medical preparations containing as active ingredient the modified antibody of the invention are useful as preventives and/or remedies for cancers, inflammation, hormone disorders and blood diseases.

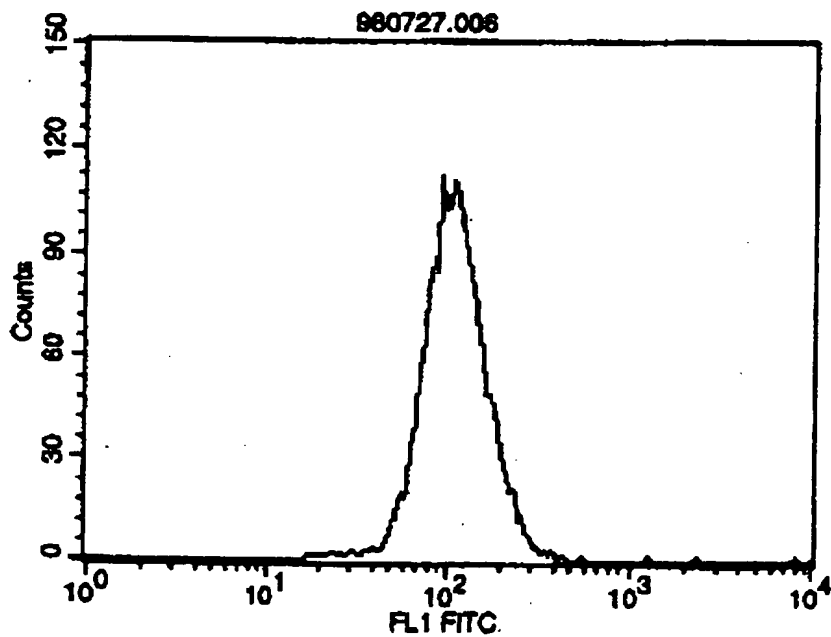
[Selected Drawings] None

【書類名】 ~~図面~~ Drawing

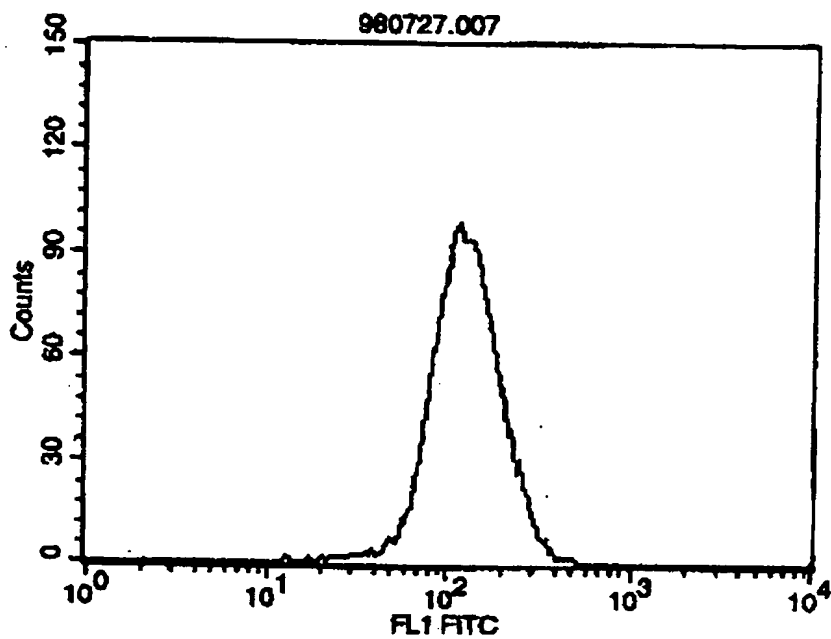
【~~図1~~】 Fig.1



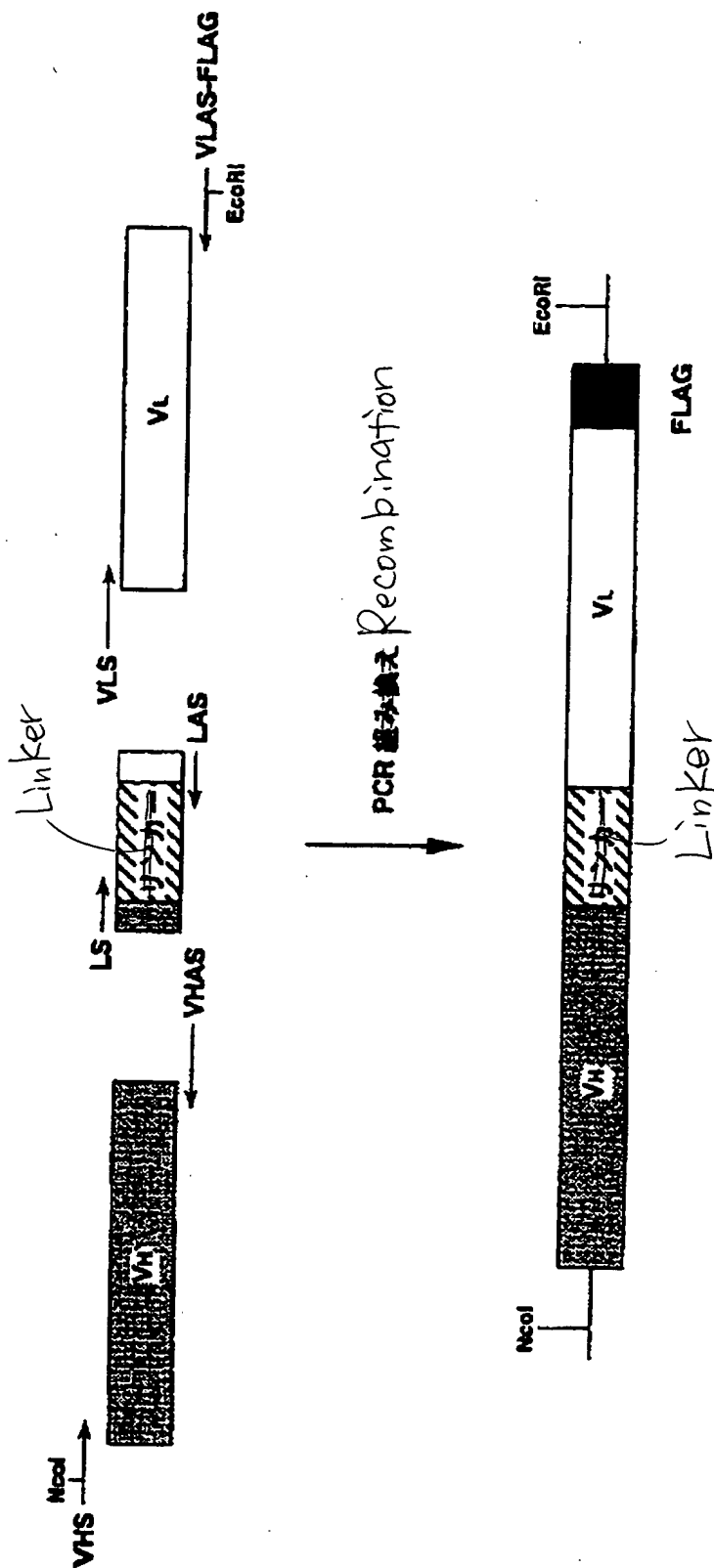
【~~図2~~】 Fig.2



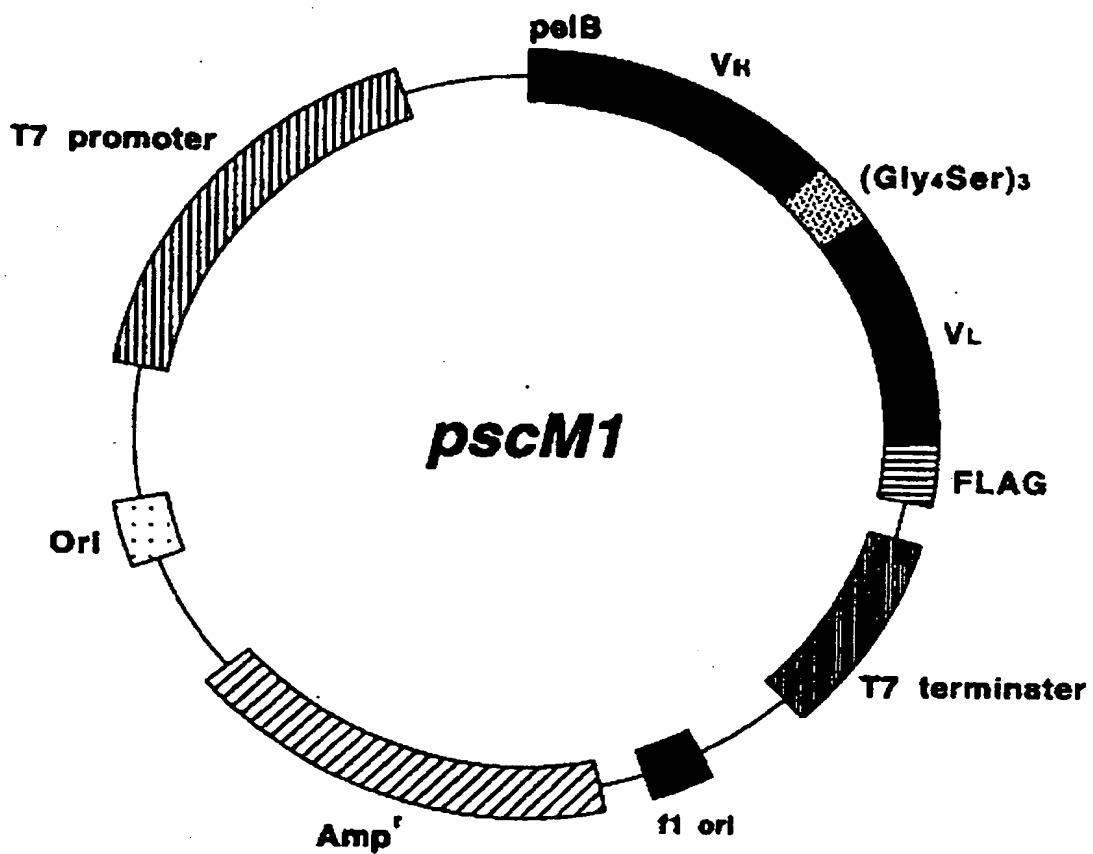
~~【図3】~~ Fig.3



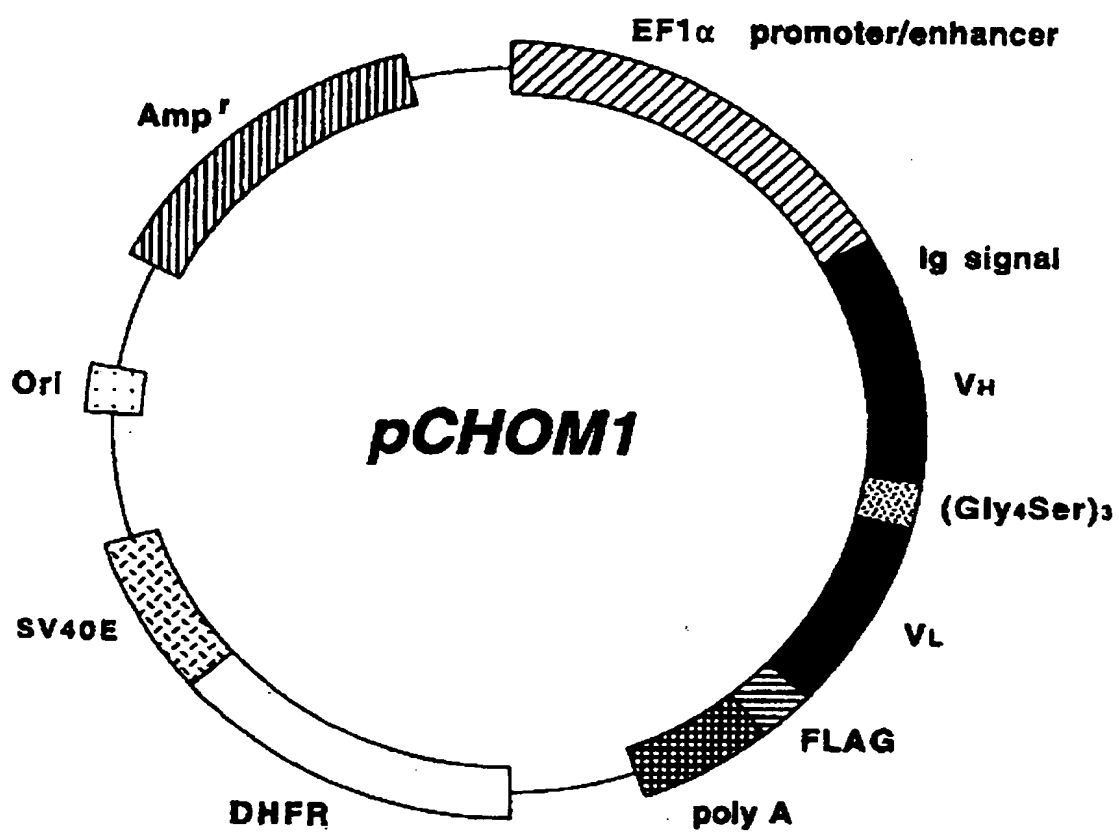
【図4】Fig.4



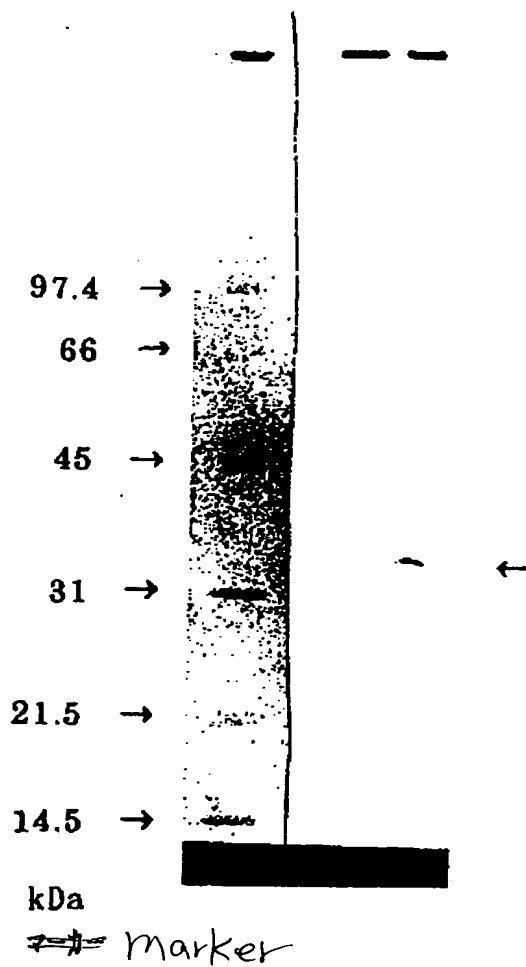
【図5】 Fig. 5.



【図6】Fig.6

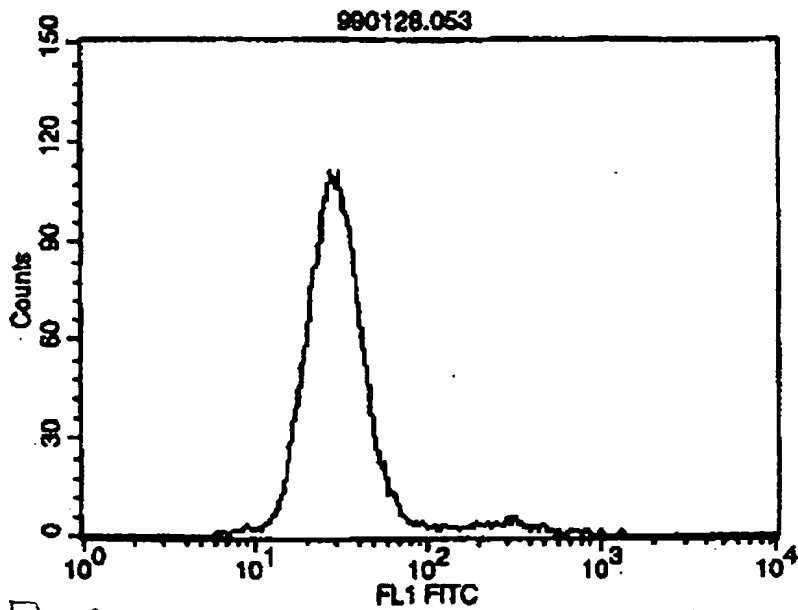


【~~図~~7】
Fig.



【~~図~~8】

(Fig.)



(Fig.)

【~~図~~9】

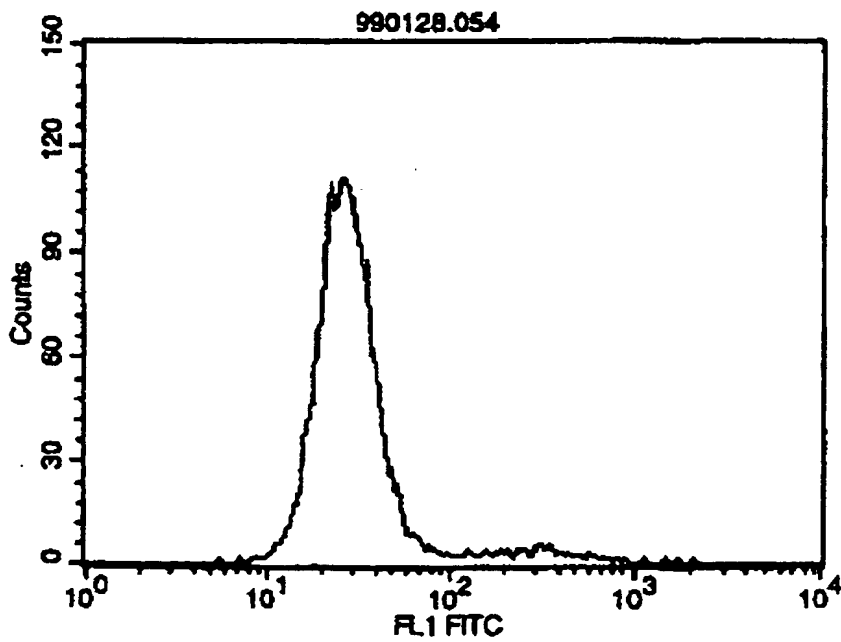


Fig. **【図10】**

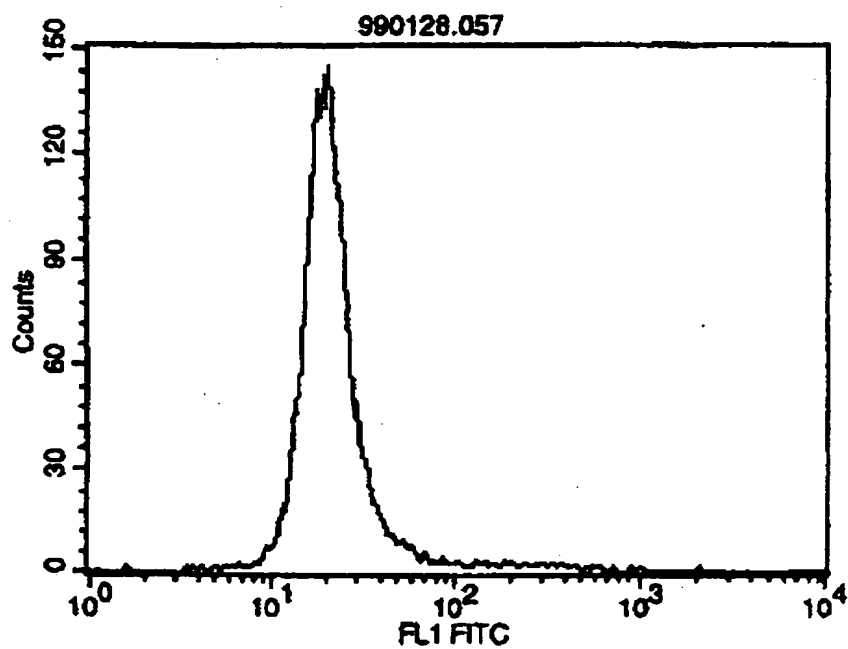
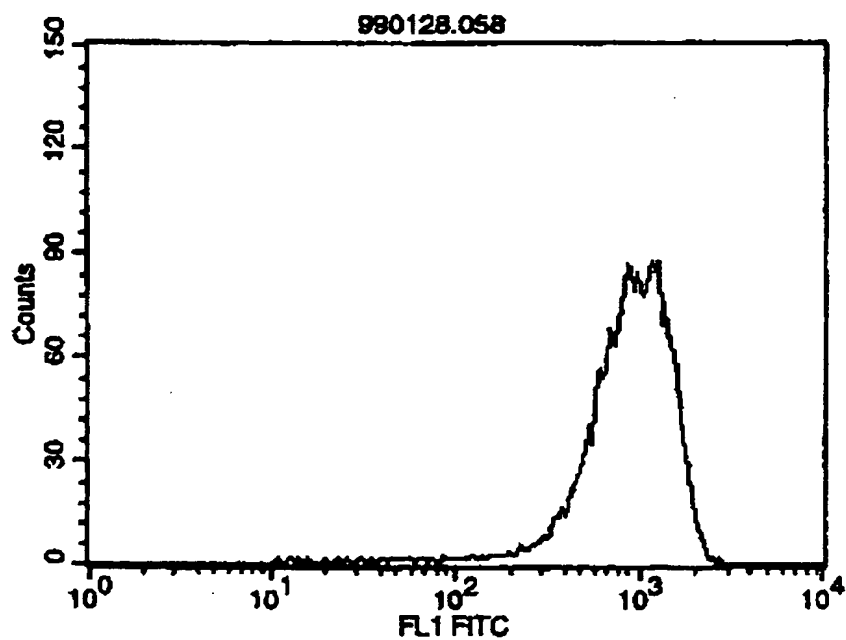


Fig. **【図11】**



【図12】

Fig.

Competitive ELISA

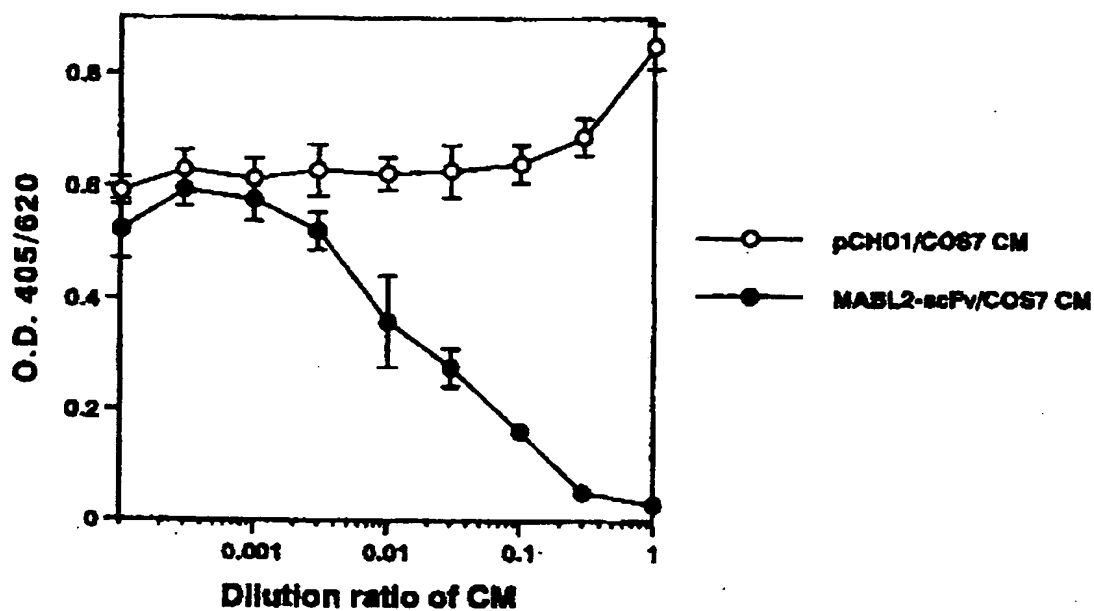
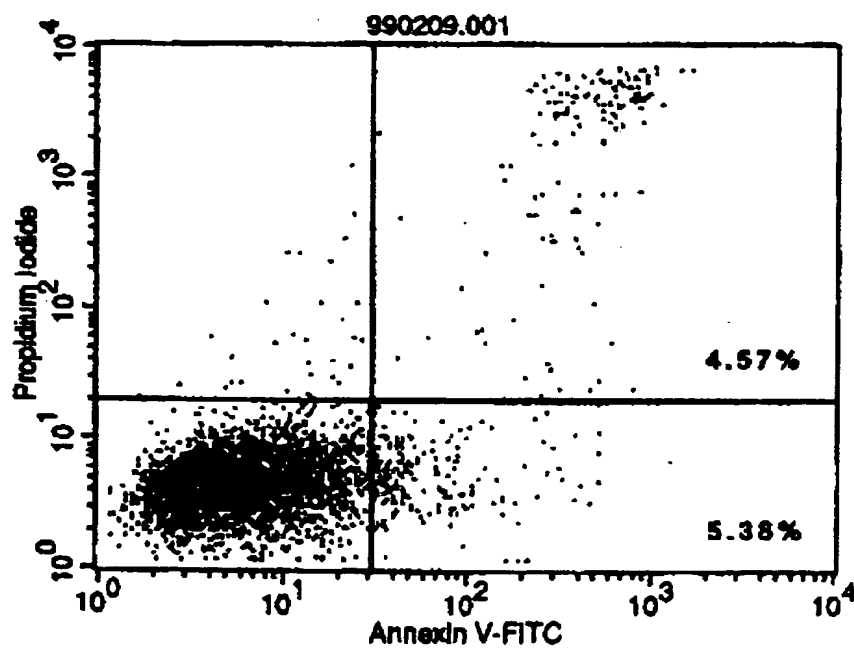


Fig.

【図13】



【図14】
Fig.

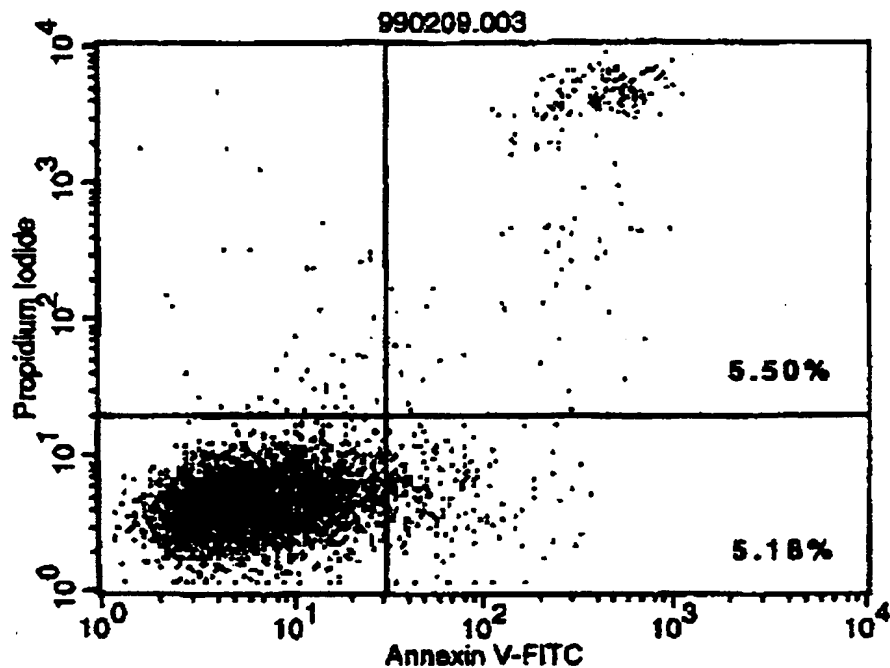
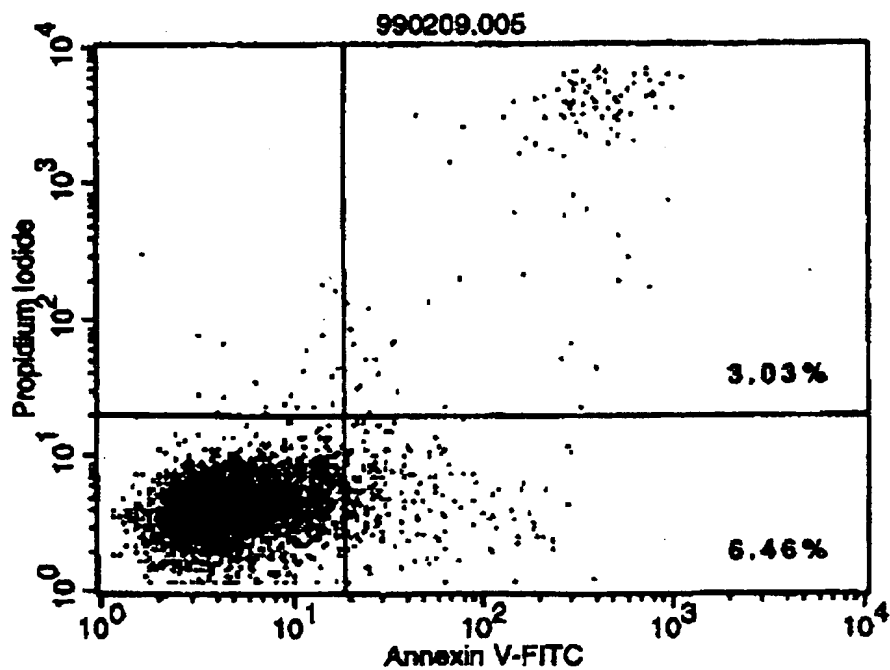


Fig.
【図15】



【図16】
Fig.

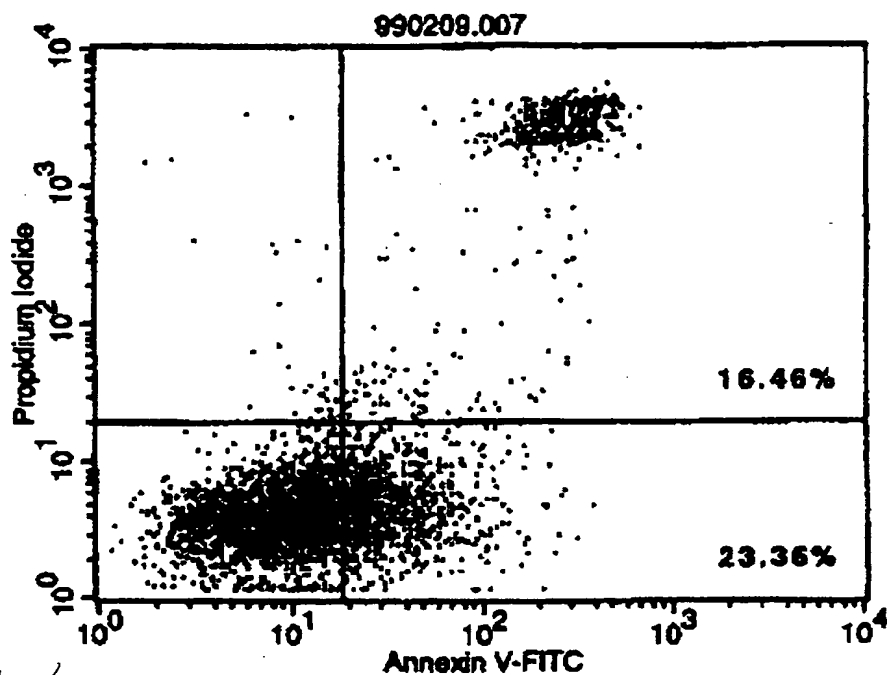
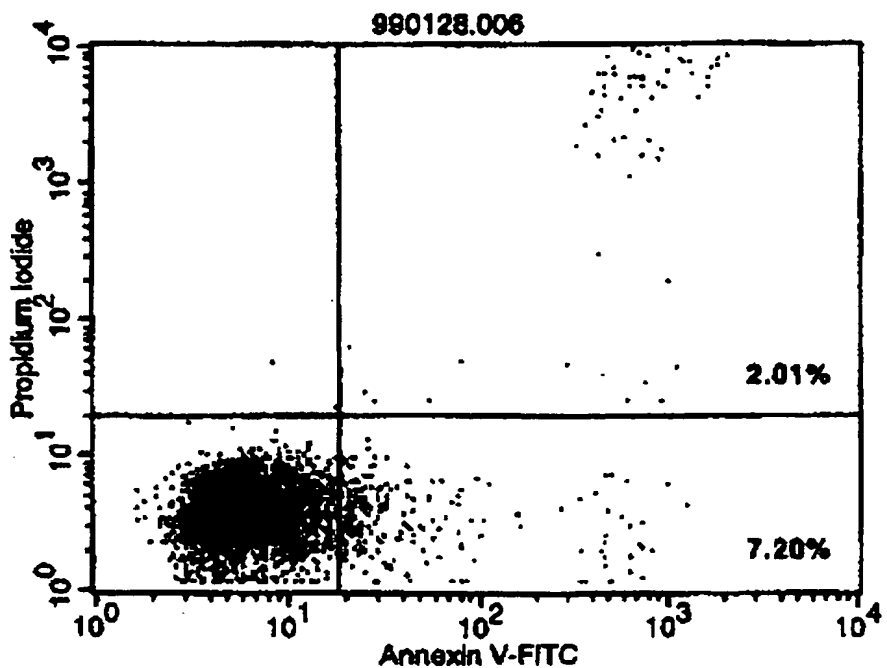
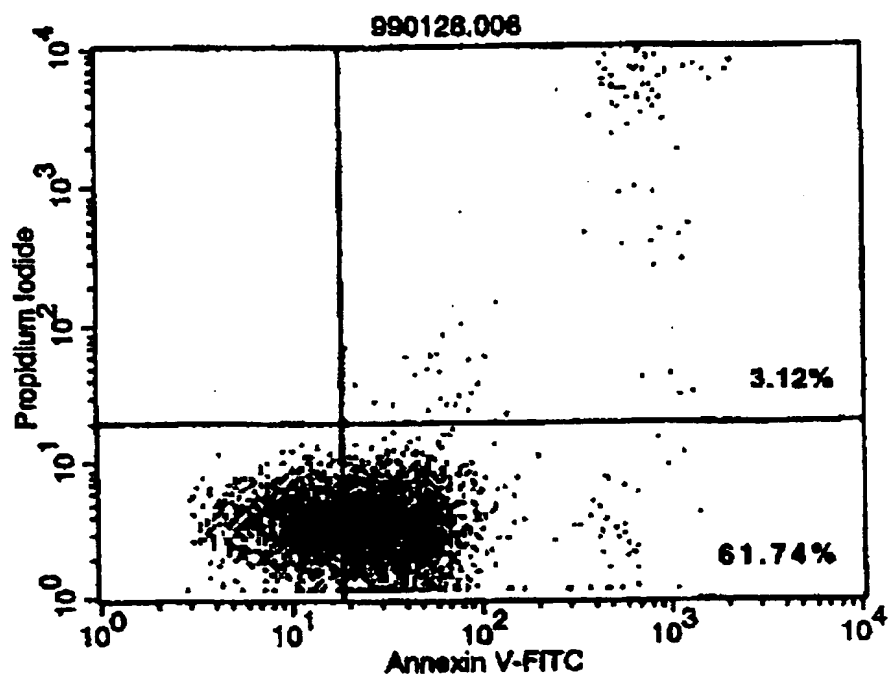


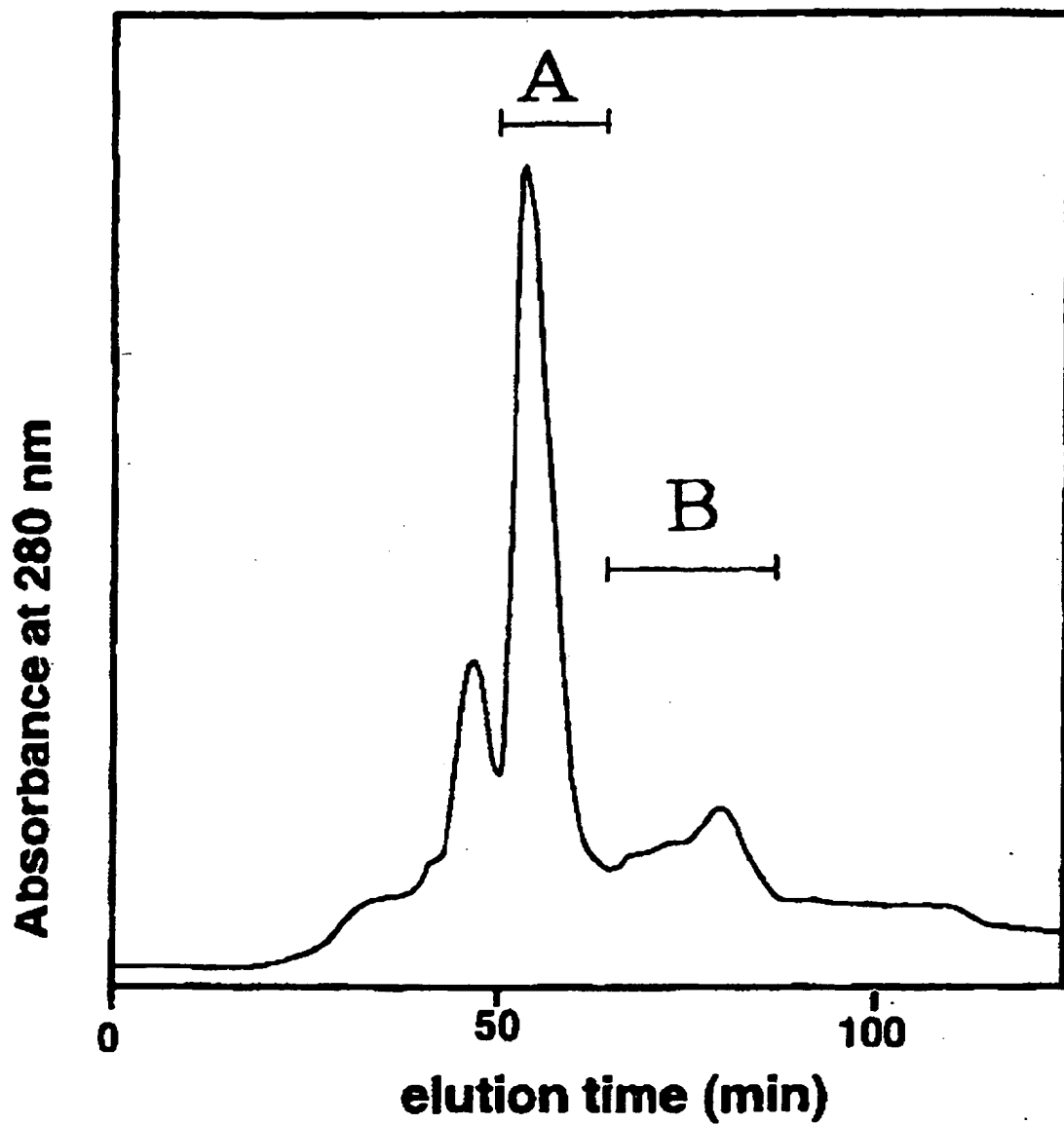
Fig.
【図17】



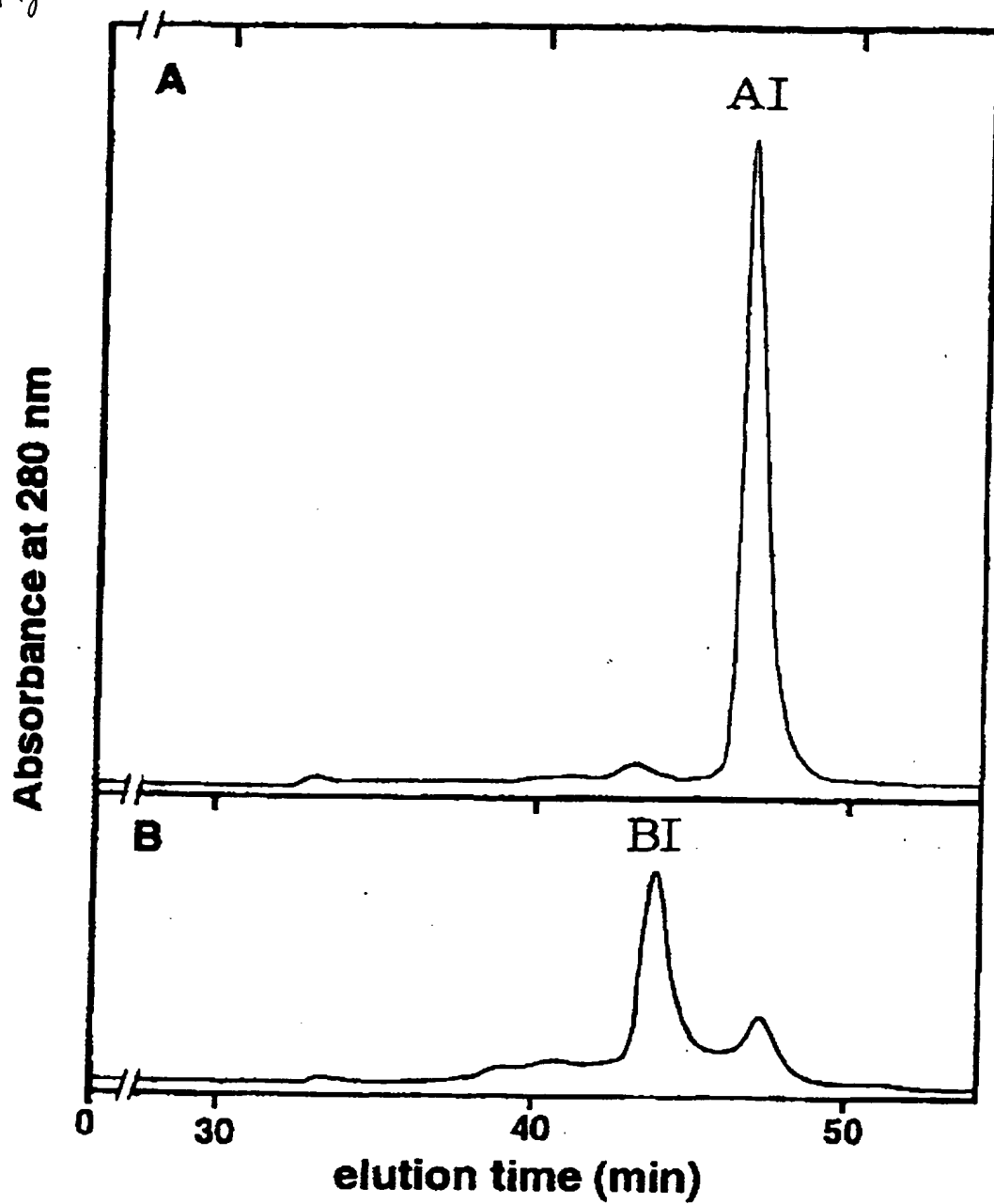
【図18】
Fig.



【図19】
F-8

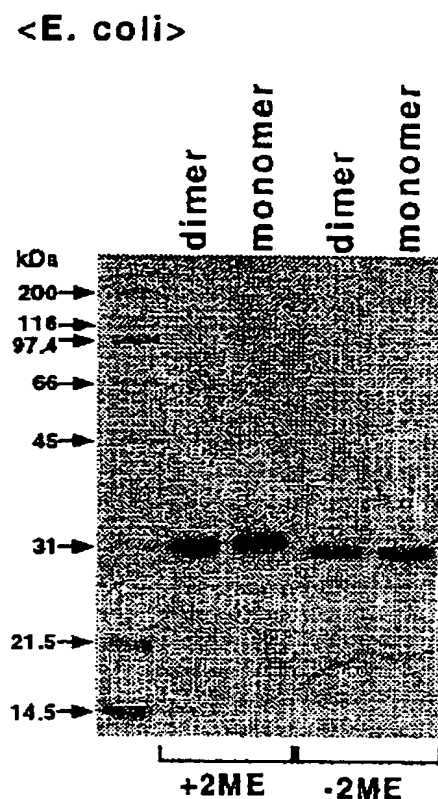
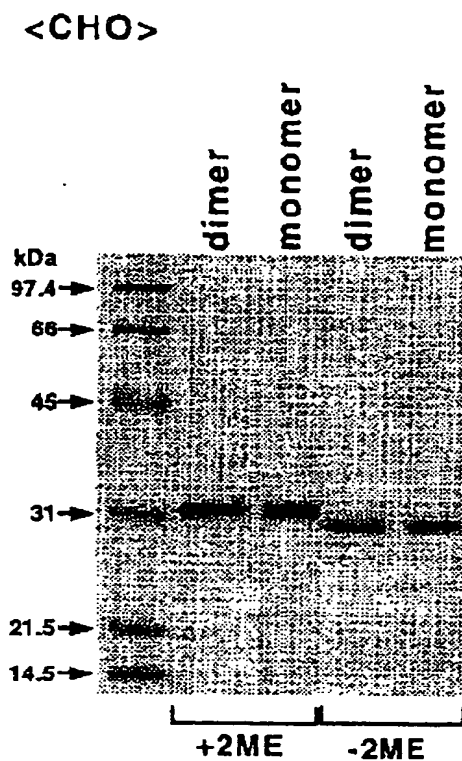


【~~図~~20】
(Fig.)



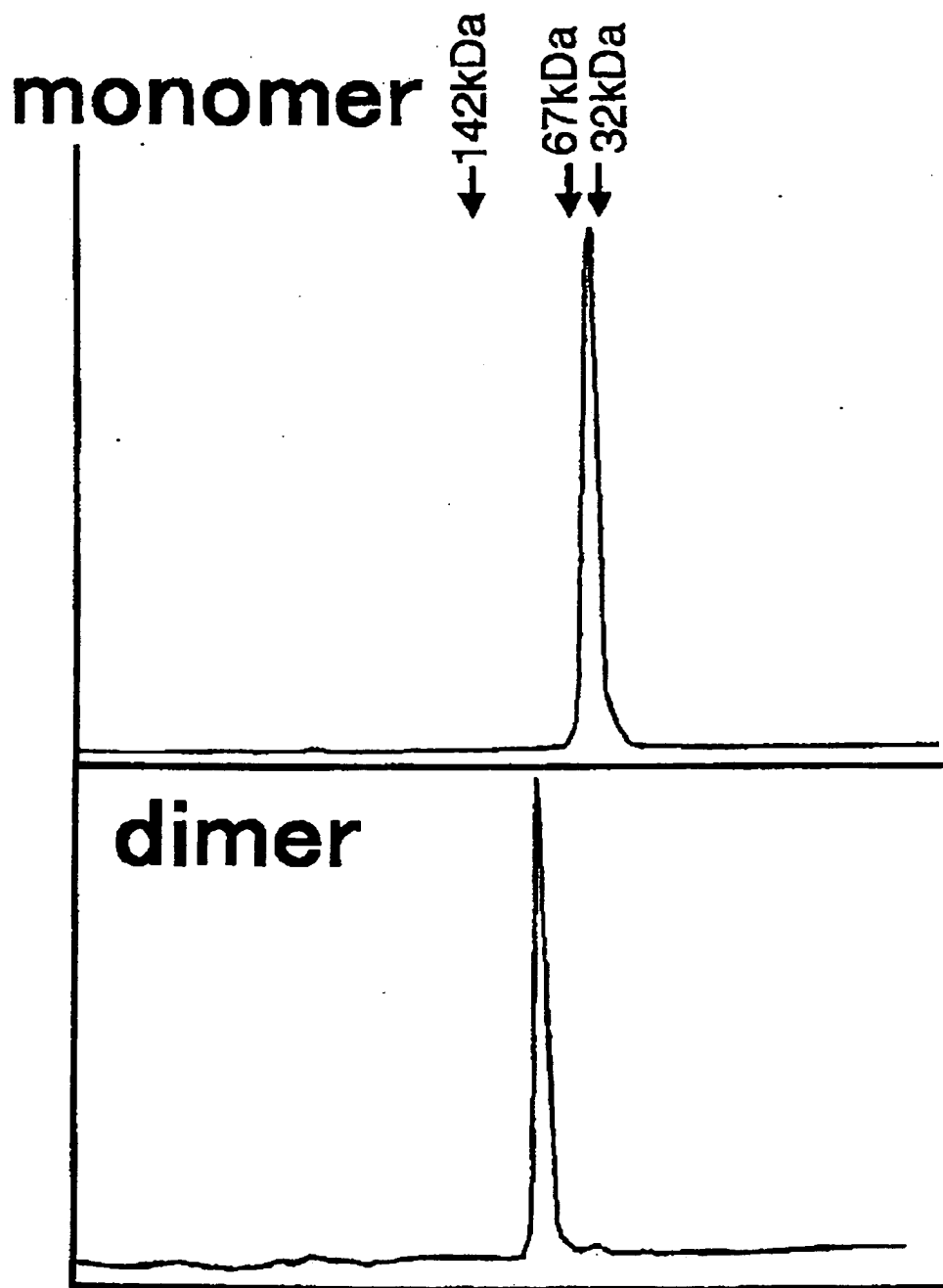
【図21】
Fig.

SDS-PAGE analysis of MABL2-scFv

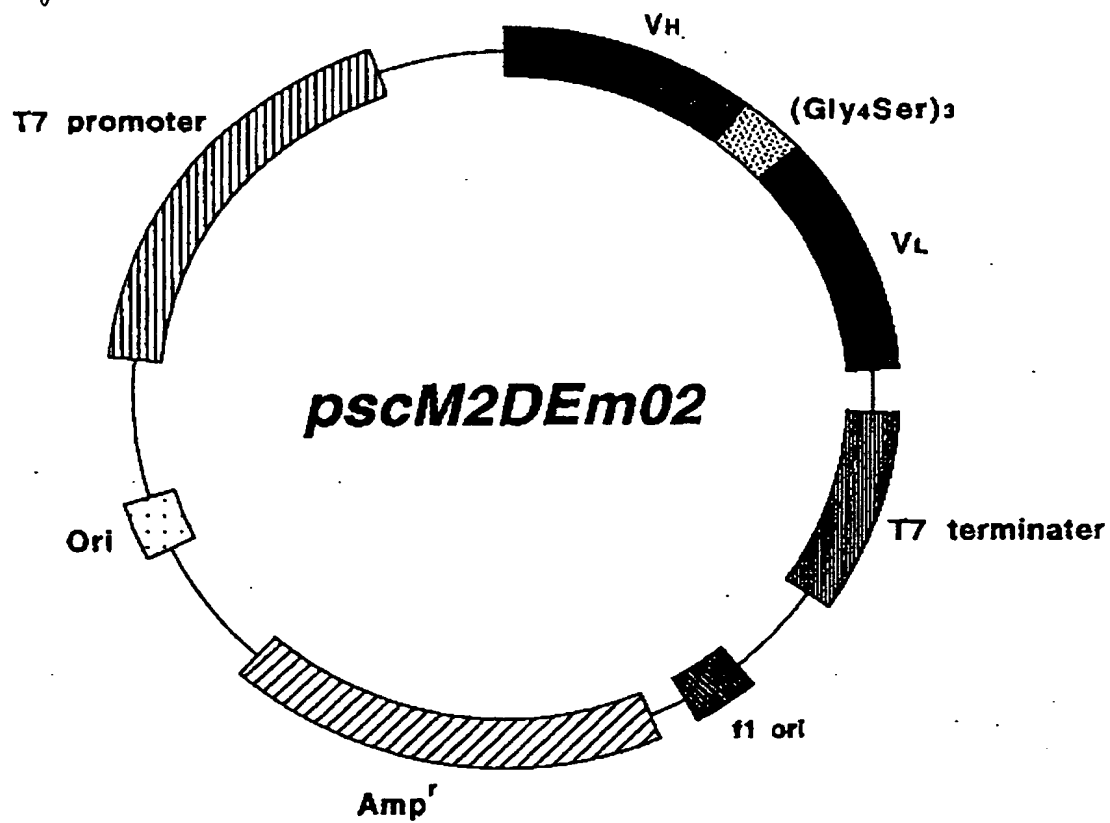


【~~図~~ 22】
(Fig.)

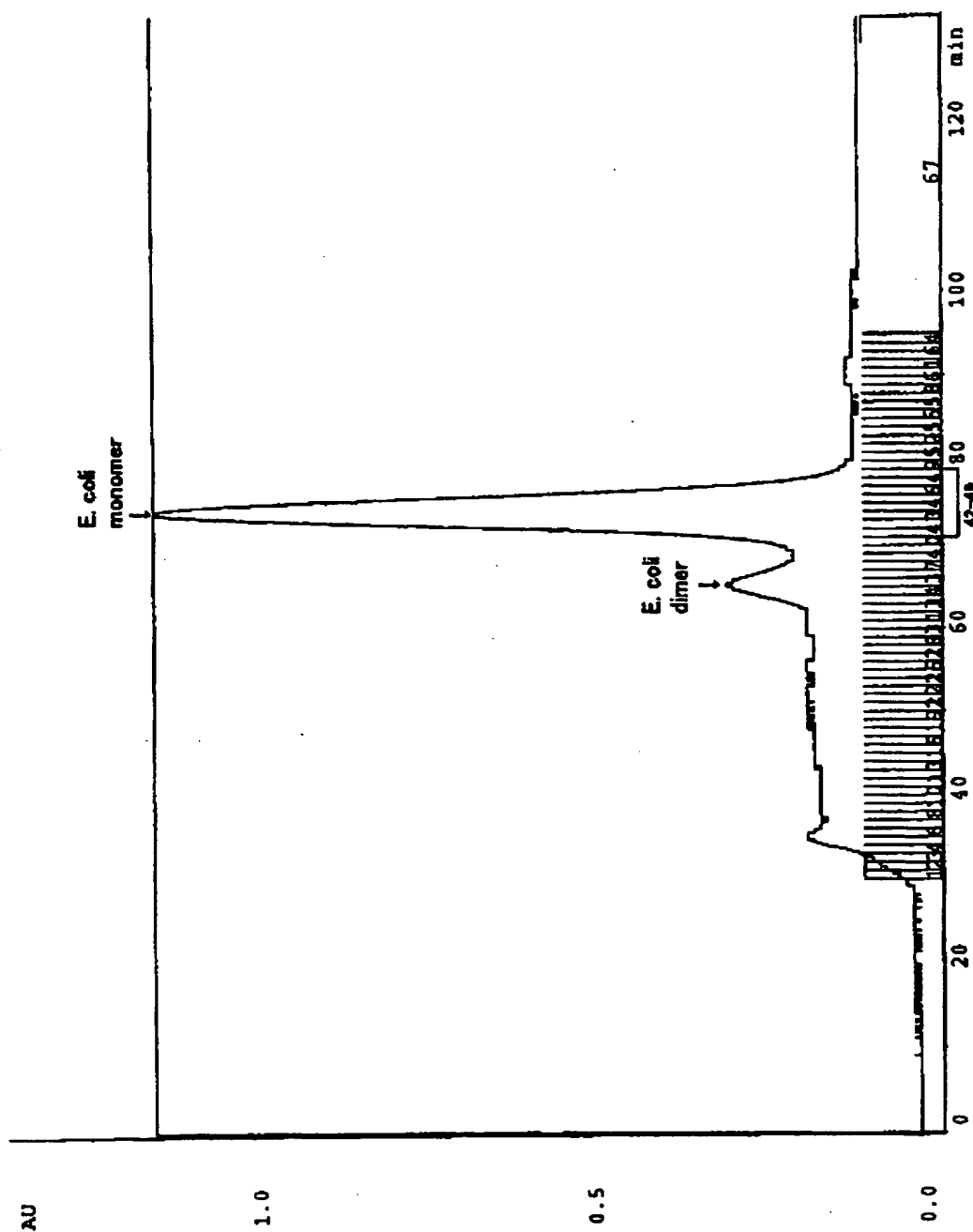
TSK gel G3000SW
20 mM Acetate buffer, 0.15 M NaCl, pH 6.0



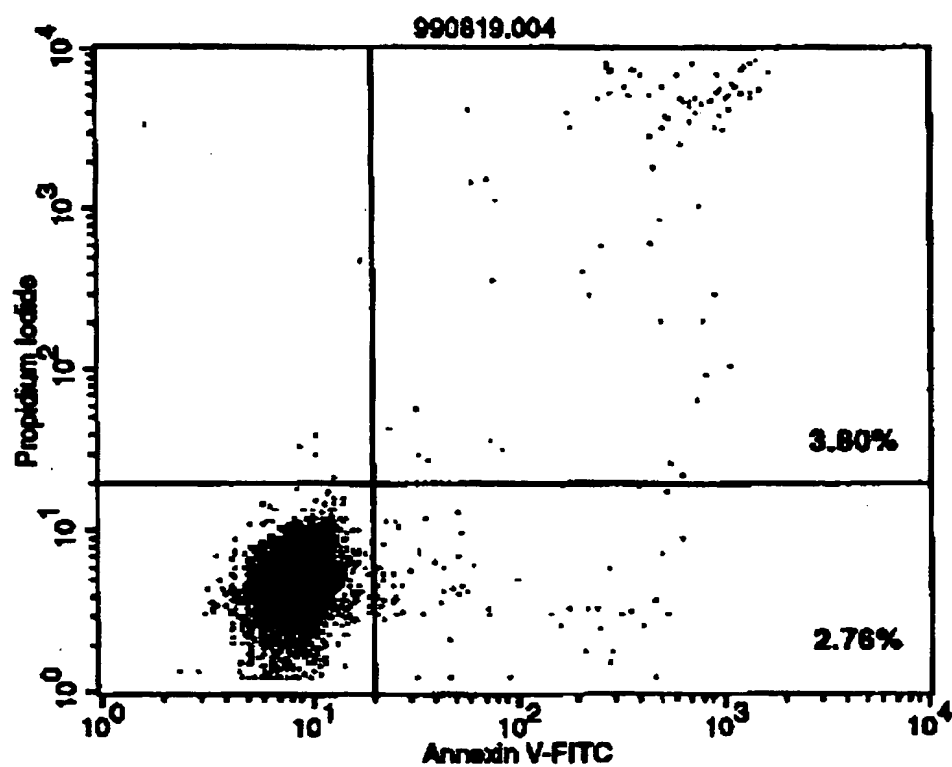
【~~図~~23】
Fig.



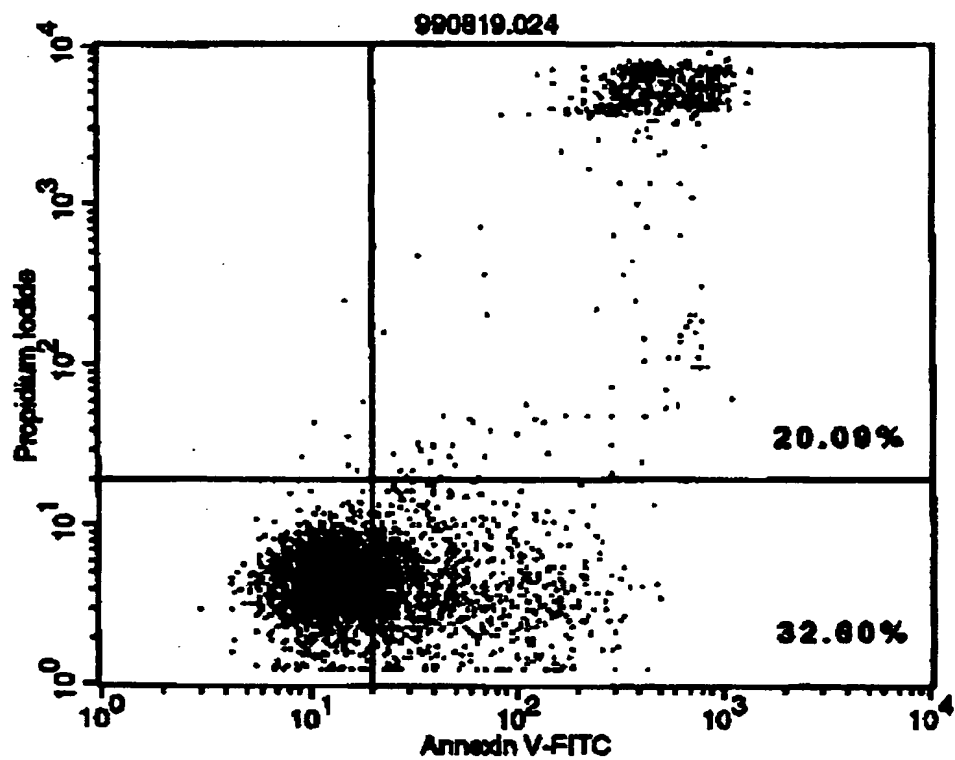
【図24】
(Fig.)



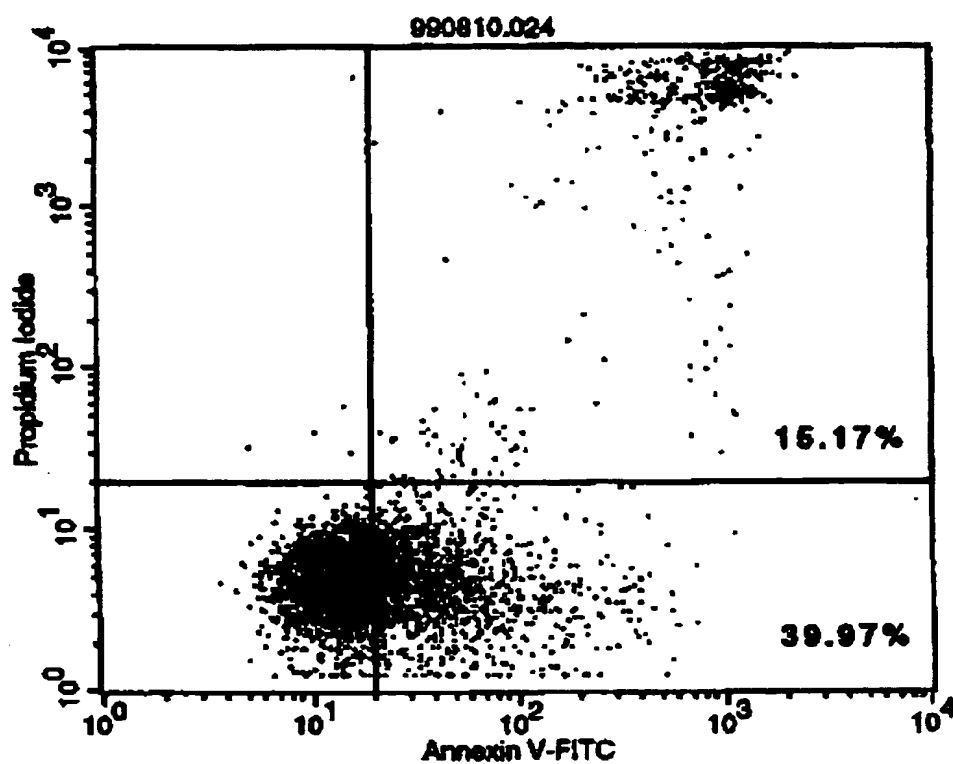
【~~図~~25】
(Fig.)



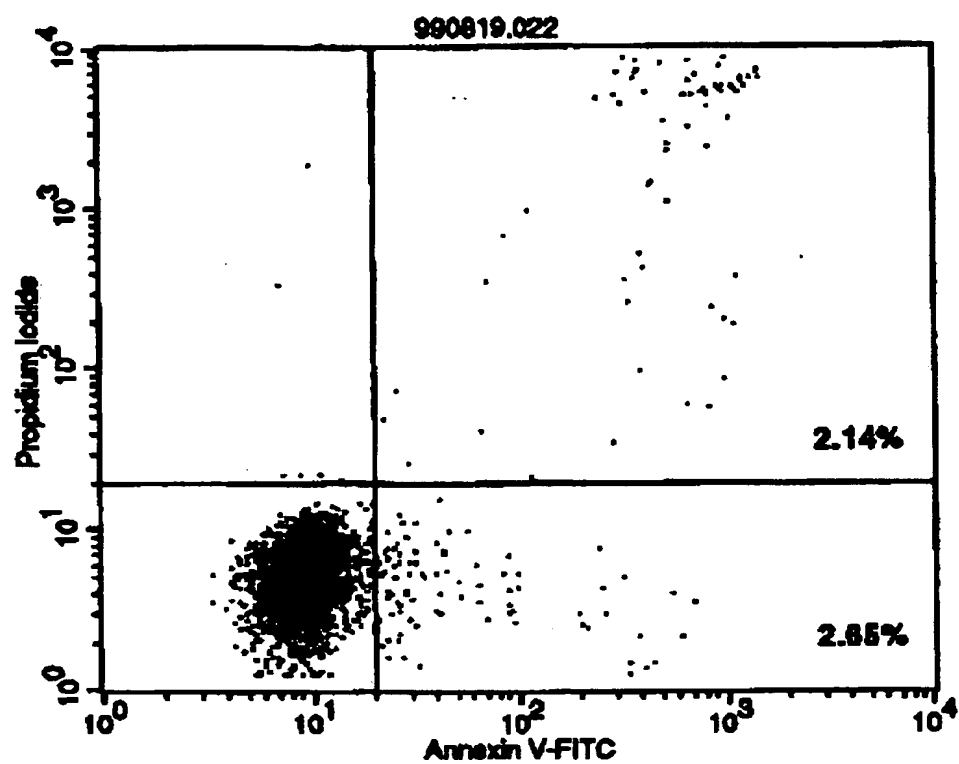
【~~図~~ 26】
F.g.



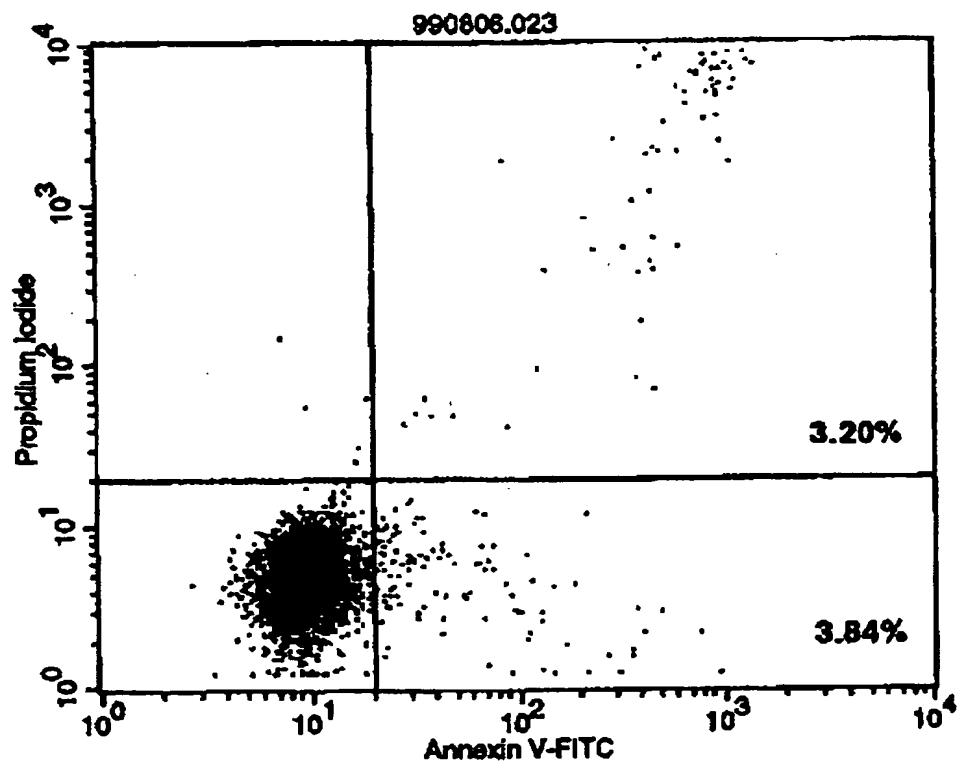
【図27】
Fig.



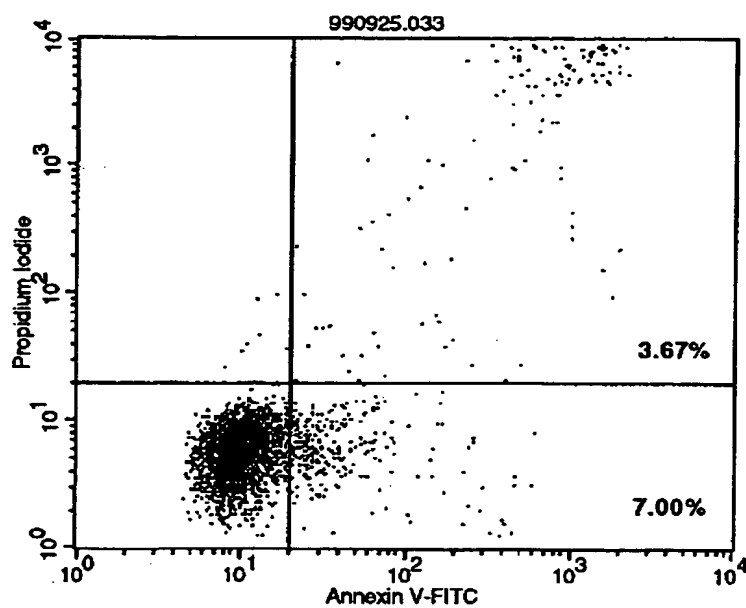
【図28】
Fig.



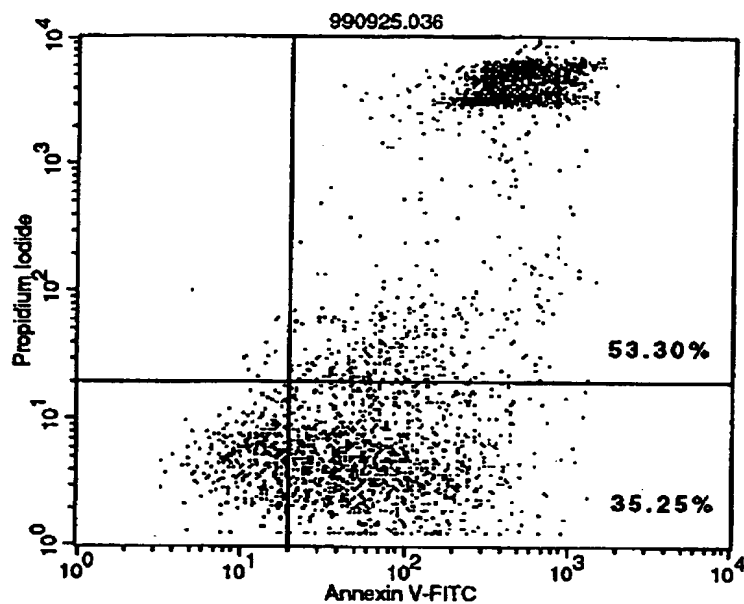
【~~図~~29】
(Fig.)



(Fig.)
【~~図~~30】

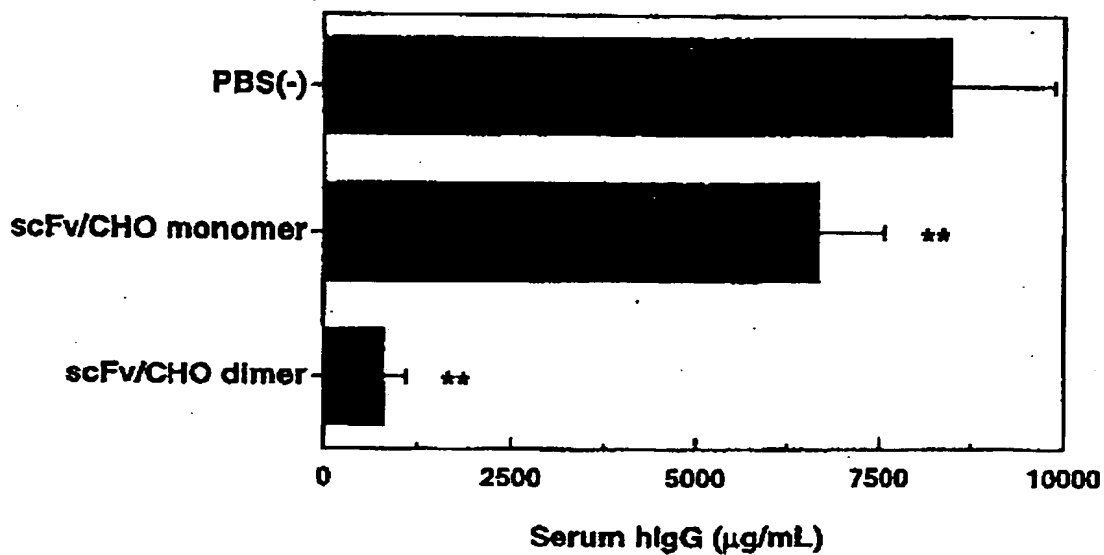


【図31】
(Fig.)



(Fig.)
【図32】

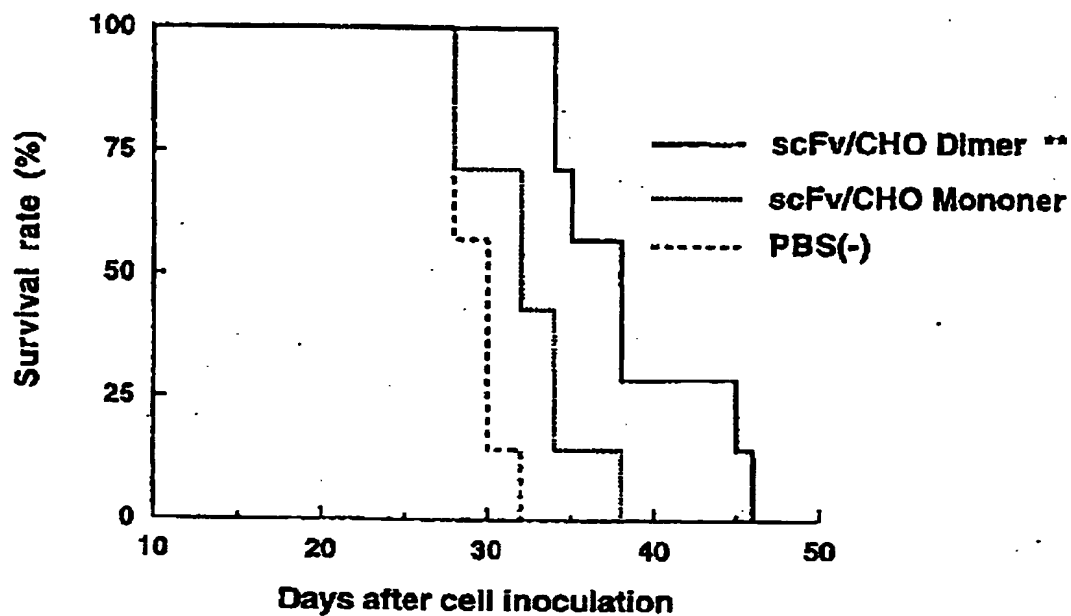
**Effect of MABL-2 (scFv) on serum hlgG
in KPM2 i.v. SCID mice**



** : $p < 0.01$

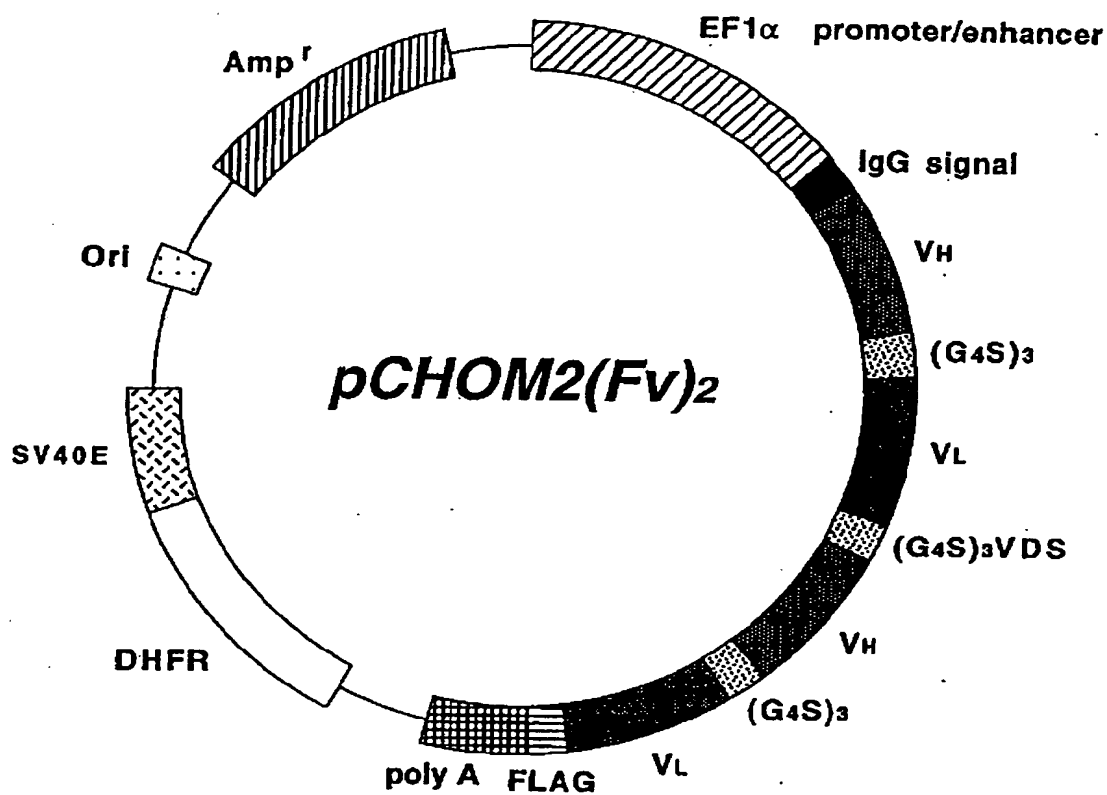
【図33】
(F.g.)

**Effect of MABL-2 (scFv) on survival
of KPMM2 i.v. SCID mice**



** ; $P < 0.01$ by t-test

【図34】
Fig.



【図35】
(Frq.)

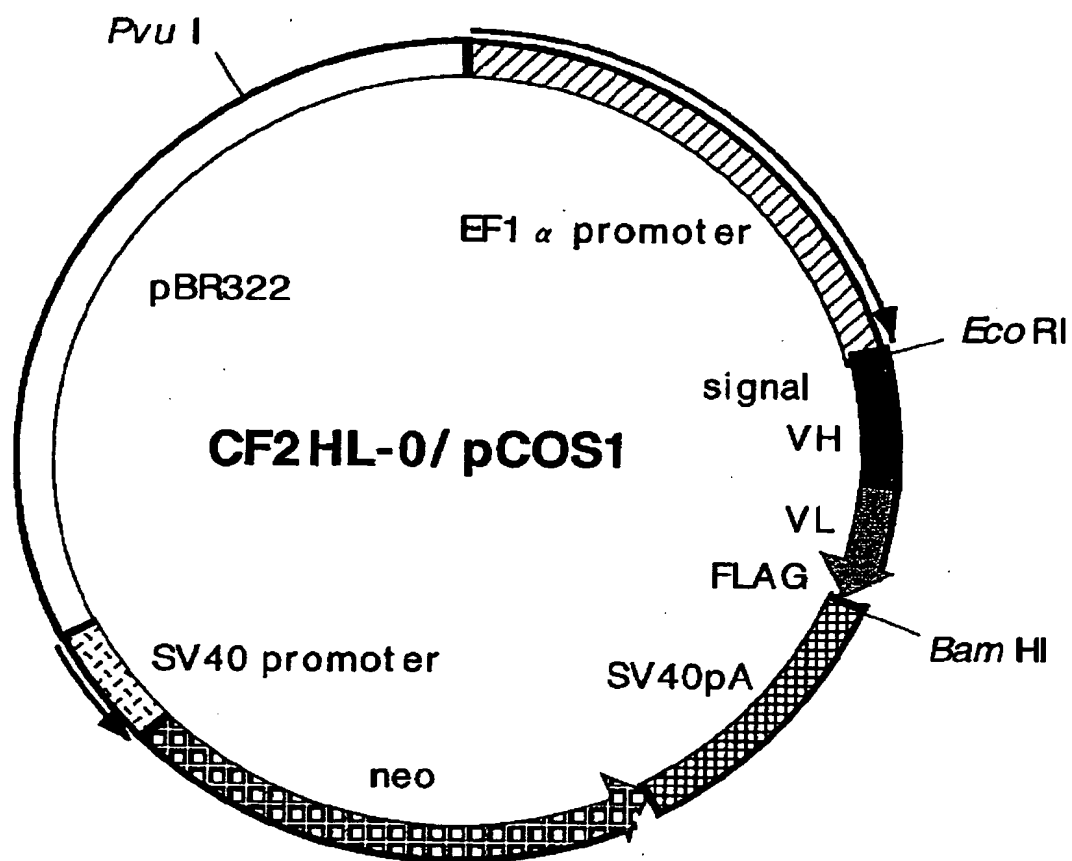
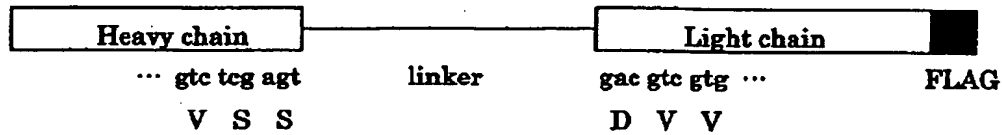


Fig. 【図36】

HL-type polypeptide and amino sequences of peptide linkers
 <HLタイプのリンカー塩基配列とアミノ酸配列>



Plasmid	Number of linker amino acid	linker	linker
CF2HL-0/pCOS1	0	gtc tcg agt V S S	gac gtc gtg D V V
CF2HL-3/pCOS1	3	gtc tcg agt ggt ggt tcc V S S G G S	gac gtc gtg D V V
CF2HL-4/pCOS1	4	gtc tcg agt ggt ggt ggt tcc V S S G G G S	gac gtc gtg D V V
CF2HL-5/pCOS1	5	gtc tcg agt ggt ggt ggt ggt tcc V S S G G G G S	gac gtc gtg D V V
CF2HL-6/pCOS1	6	gtc tcg agt gt ggt ggt ggt ggt tcc V S S G G G G G S	gac gtc gtg D V V
CF2HL-7/pCOS1	7	gtc tcg agt ggt ggt ggt ggt ggt ggt tcc V S S G G G G G G S	gac gtc gtg D V V

【図37】

Fig.

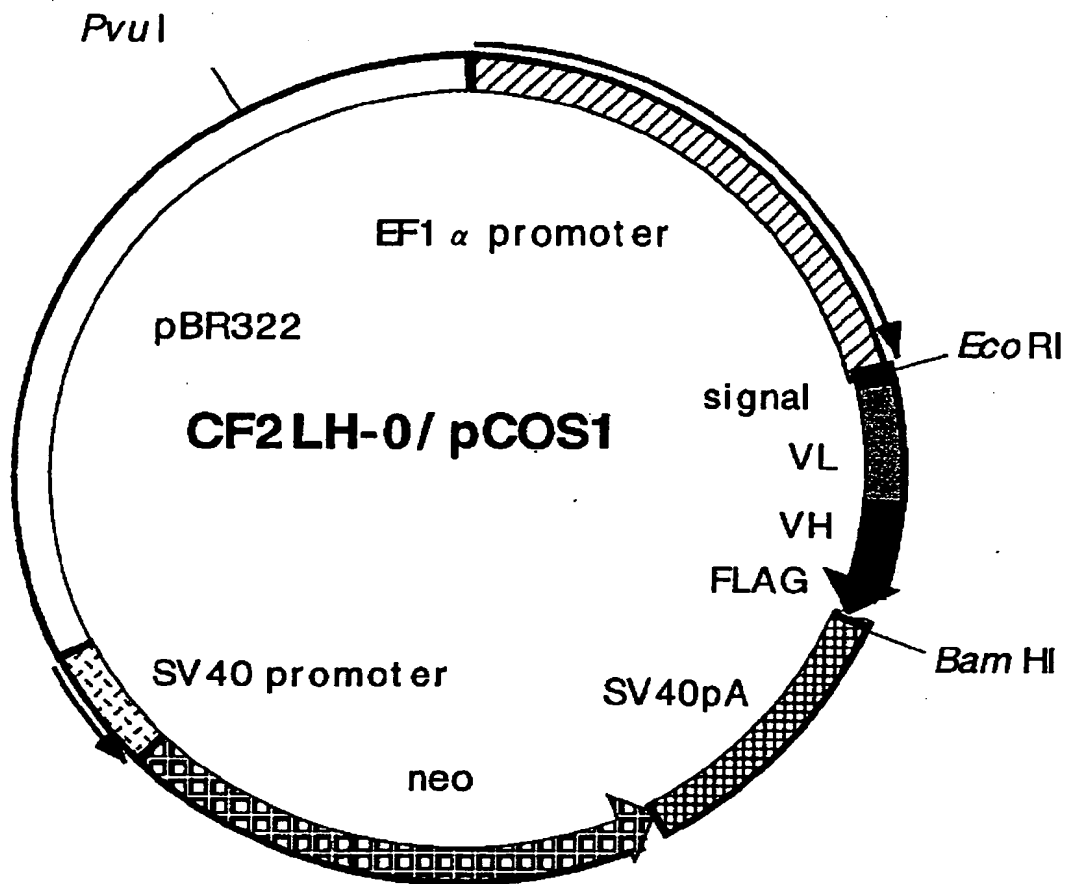
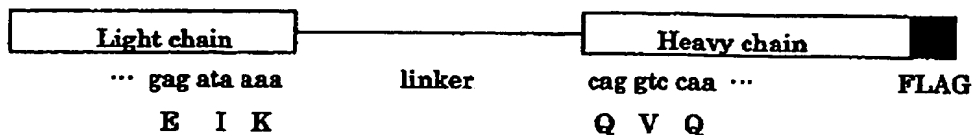
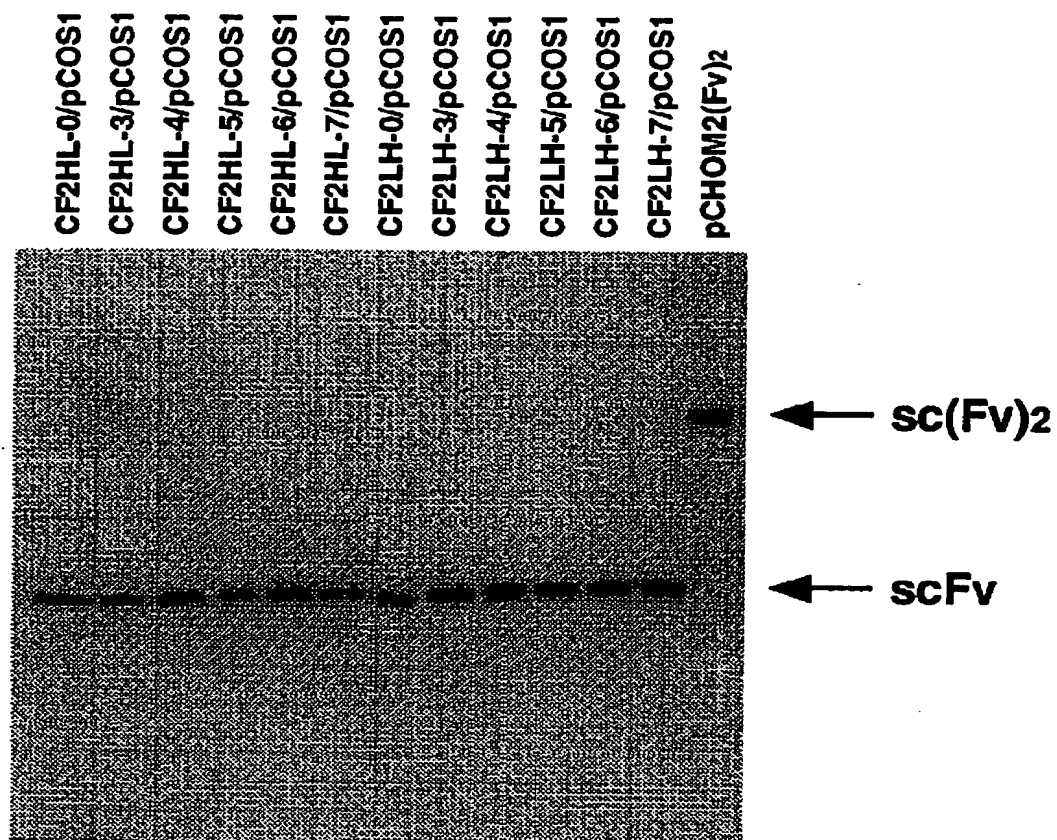


Fig. 【図38】

LH-type polypeptide and amino acid sequences of peptide linkers
＜LHタイプのリンカー塩基配列とアミノ酸配列＞

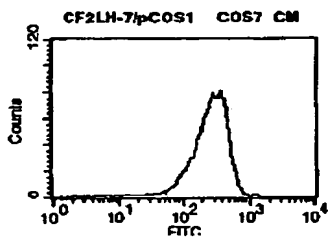
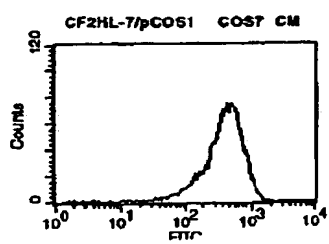
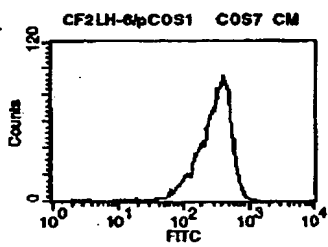
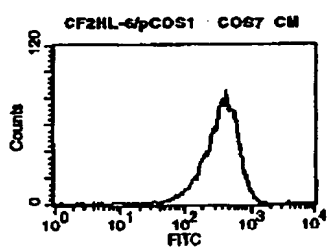
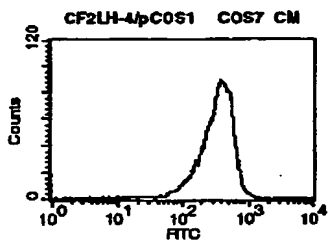
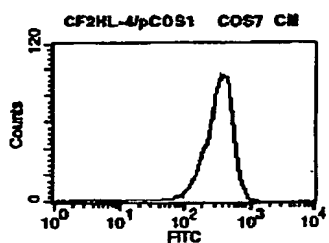
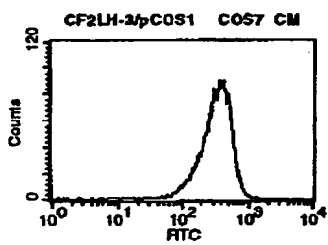
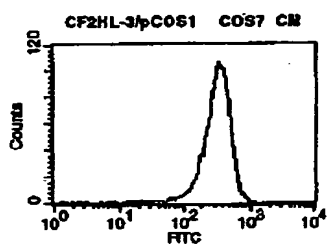
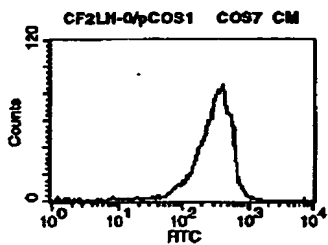
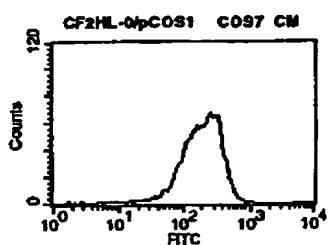
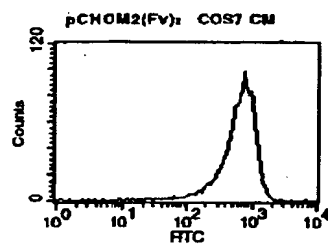
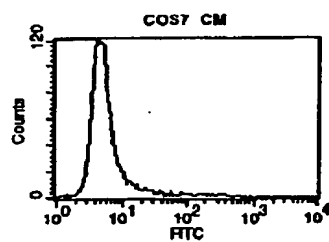
Plasmid	Number of linker amino acid	linker	linker
CF2LH-0/pCOS1	0	gag ata aaa E I K	cag gtc caa Q V Q
CF2LH-3/pCOS1	3	gag ata aaa tcc gga ggc E I K S G G	cag gtc caa Q V Q
CF2LH-4/pCOS1	4	gag ata aaa tcc gga ggt ggc E I K S G G G	cag gtc caa Q V Q
CF2LH-5/pCOS1	5	gag ata aaa tcc gga ggt ggt ggc E I K S G G G G	cag gtc caa Q V Q
CF2LH-6/pCOS1	6	gag ata aaa tcc gga ggt ggt ggt ggc E I K S G G G G G	cag gtc caa Q V Q
CF2LH-7/pCOS1	7	gag ata aaa tcc gga ggt ggt ggt ggt ggc E I K S G G G G G G	cag gtc caa Q V Q

【~~図~~39】
Fig.

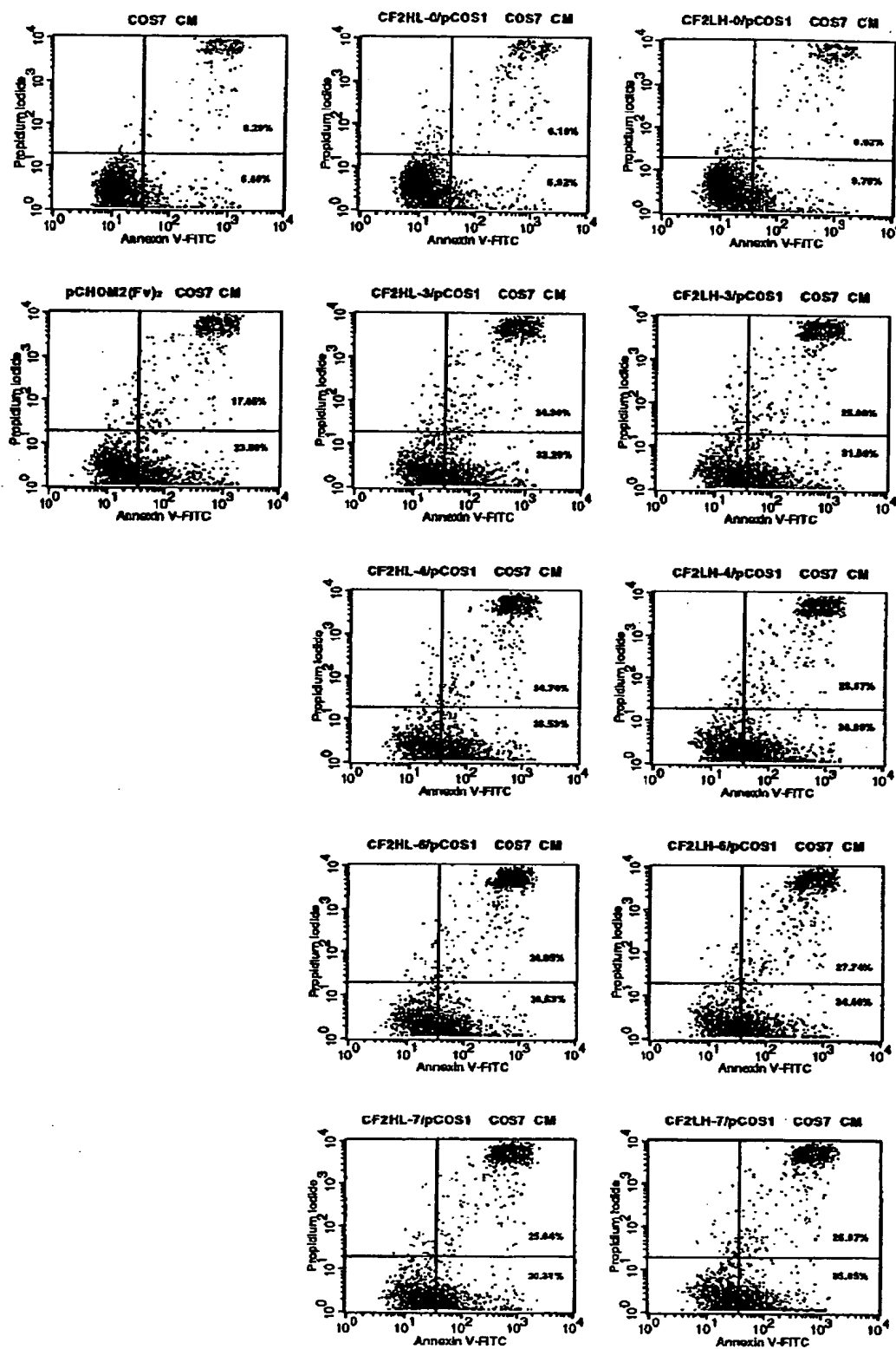


【図40】

Fig.

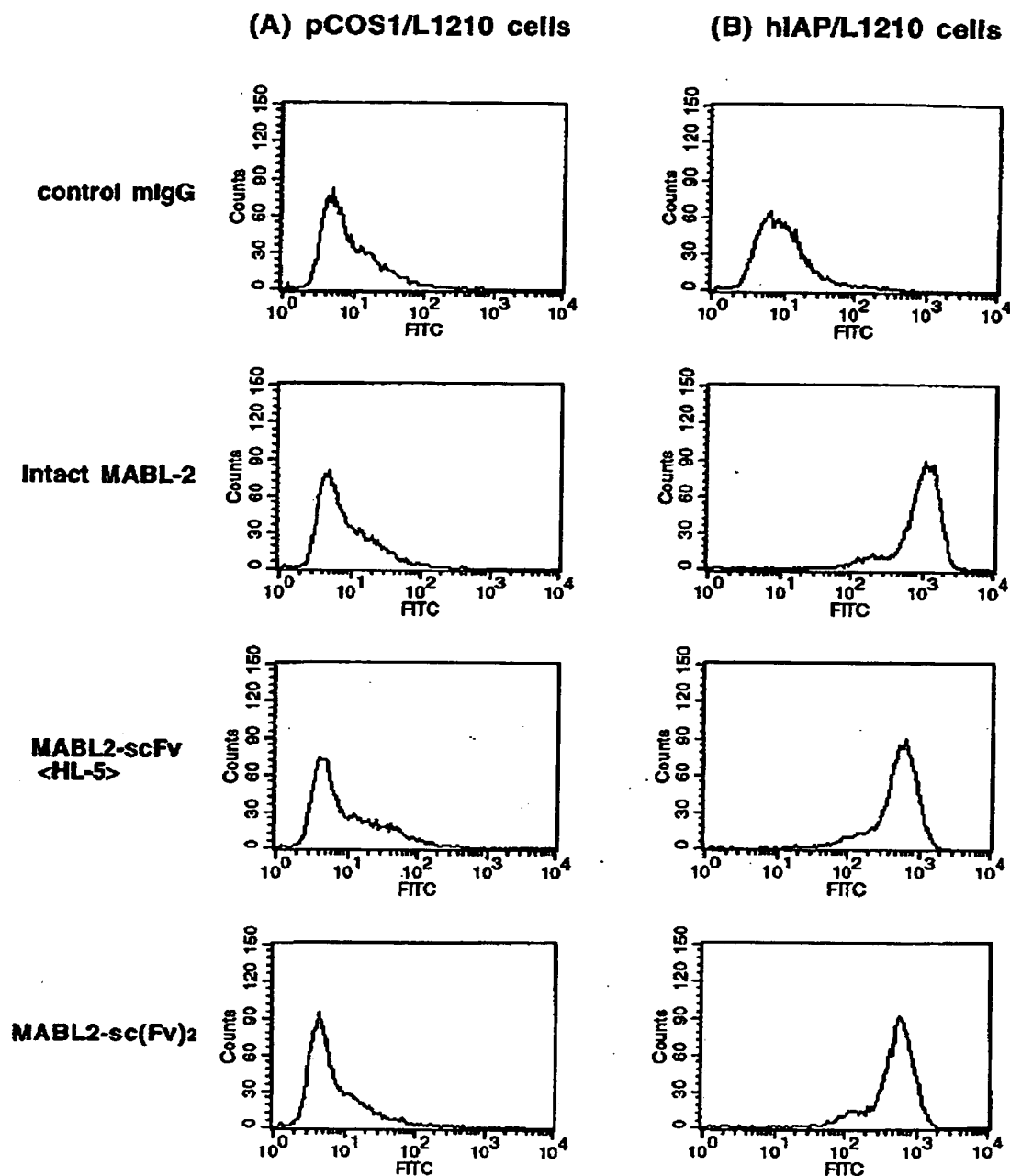


【図41】
Fig.



【図42】

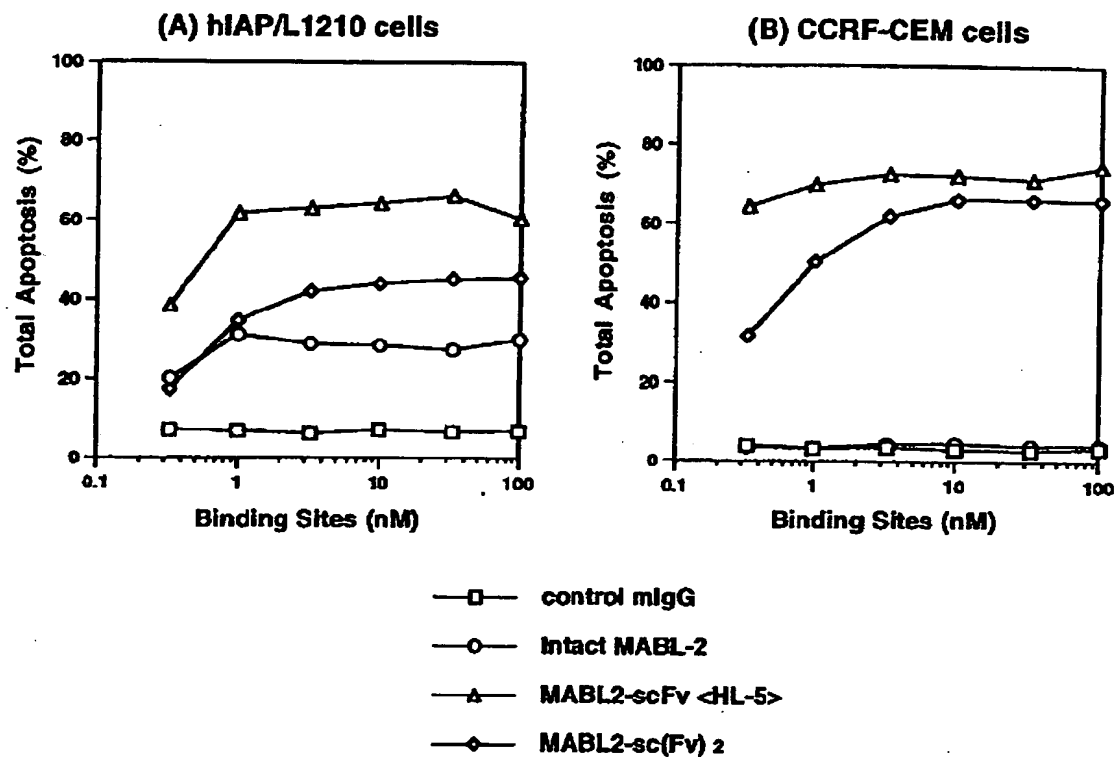
Reactivity of MABL-2 derivatives with pCOS1/L1210 (A) and hIAP/L1210 (B) cells



【図43】

(Fig.)

**Apoptosis on hIAP/L1210 (A) and CCRF-CEM (B) cells
by Annexin V-staining**



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